## SEQUENCE LISTING

<110>	LIPSCOMBE, Diane BELL, Thomas THALER, Christopher CASTIGLIONI, Andrew	
<120>	NOCICEPTIVE NEURON-SPECIFIC CALCIUM CHANNEL ISOFORM AND USES THEREOF	
<130>	B0877.70026US00	
	US 60/443,474 2003-01-29	
<160>	47	
<170>	PatentIn version 3.2	
<210><211><212><213>		
<220>	·	
	PCR primer	
<400> gctgcgt	1 Egtt geeggattea ttat	24
	24	
<220> <223>	PCR primer	
	2 cgaa ccaggegett gtag	24
<210><211><212><212><213>	3 24 DNA Artificial	
<220> <223>	PCR primer	
<400> ctgaata	3 acga cccagctgeg tgtg	24
<210><211>	4 24	

1

	DNA Artificial	
<220> <223>	PCR primer	
<400>	4	
ccaggc	gett gtatgeaact egag	24
<210>	5	
	22	
<212>		
<213>	Artificial	
<220>		
<223>	PCR primer	
-400-		
<400>	ttot ggagoottag ot	
cgcagg	ggageetag et	22
010		
<210> <211>		
<212>		
	Artificial	
<220>		
	PCR primer	
12237	rek primer	
<400>		
ggccat	tgct gtggacaacc tt	22
<210>	7	
<211>	28	
<212>		
<213>	Artificial	
<220>		
<223>	PCR primer	
<400>	7	
	ggac ttcattgttg tcagtgga	28
		20
<210>	8	
<211>	28	
<212>	DNA	
<213>	Artificial	
<220>		
<223>	PCR primer	
-100-	O.	
<400>	8 gact ctcagagact tgatggta	
cegeage	gaet eccayayact tyatgyta	28

.

```
<210> 9
<211> 25
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 9
cagagatgcc tggaacgtct ttgac
                                                                     25
<210> 10
<211> 25
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 10
ataacaagat gcggatggtg tagcc
                                                                     25
<210> 11
<211> 30
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 11
agctgcgtgt tgccggattc attataagga
                                                                     30
<210> 12
<211> 30
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 12
gcgcttgtag gccaacctac gagggcagtt
                                                                     30
<210> 13
<211> 30
<212> DNA
<213> Artificial
<220>
<223> PCR primer
```

Ţ

```
<400> 13
agctgcgtgt gggcgcatca gttacaatga
                                                                     30
<210> 14
<211> 30
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 14
gcgcttgtat gcaactcgag ccgggcattt
                                                                     30
<210> 15
<211> 26
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 15
tcaattccca cacacctccc agttcc
                                                                     26
<210> 16
<211> 26
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 16
                                                                     26
tttgggtggt ctgcttaggg aaccag
<210> 17
<211> 24
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 17
                                                                     24
gcgctggttg taagggtcag attg
<210> 18
<211> 24
<212> DNA
<213> Artificial
```

```
<220>
<223> PCR primer
<400> 18
agccaggcaa caattgcaga aatc
                                                                    24
<210> 19
<211> 22
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 19
cagcaccagc atcaccccat tt
                                                                    22
<210>
      20
<211> 24
<212> DNA
<213> Artificial
<220>
<223> PCR primer
<400> 20
caagatggtg aaggtcggtg tgaa
                                                                    24
<210> 21
<211> 135
<212> DNA
<213> Artificial
<220>
<223> Competitive template for exon 37a
<400> 21
agetgegtgt tgeeggatte attataagga taagegtget ggaetetete gaegeaaate
gaggcccaat tggcaaatac agttgcagaa caggaggagc cacacaactg ccctcgtagg
                                                                   120
ttggcctaca agcgc
                                                                   135
<210> 22
<211> 135
<212> DNA
<213> Artificial
<220>
<223> Competitive template for exon 37b
<400> 22
agctgcgtgt gggcgcatca gttgcaatga taagcgtgct ggactctctc gacgcaaatc
```

```
qaqqcccaat tggcaaatac agttgcagaa caggaggagc cacacaaaag cccggctcga
gttgcataca agcgc
                                                                     135
<210> 23
<211> 9562
<212> DNA
<213> Rattus norvegicus
<220>
<221> exon
<222>
      (1)..(128)
<223> Exon 36, continues 5' to nt 1
<220>
<221> misc_feature
<222> (1057)..(1066)
<223> n = a, c, g or t/u
<220>
<221> exon
<222> (5212)..(5308)
<223> Exon 37a, alternatively spliced exon
<220>
<221> exon
<222> (6798)..(6894)
<223> Exon 37b, alternatively spiced exon
<220>
<221> misc_feature
<222> (8689)..(8689)
<223> n = a, c, g or t/u
<220>
<221> misc_feature
<222> (8721)..(8721)
<223> n = a, c, g or t/u
<220>
<221>
      exon
<222>
      (9454)..(9562)
<223> Exon 38, continues 3' to nt 9562
<400> 23
atg ctg aac ctc ttt gtt gct gtg atc atg gac aat ttc gaa tac ctc
                                                                      48
Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu Tyr Leu
acg cgg gat tot toe atc cta ggg ccg cac cac ctc gat gaa ttc att
                                                                      96
Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu Phe Ile
cgc gtc tgg gct gaa tac gac cca gct gcg tg gtgagtgagc tccagtgctc
```

## Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys 35 40

tetgeteete agggttgget	gccattgcct	gtggttctcc	agacagggcc	agggtgatct	208
tagacagttc catgtgcatg	tccctaggtg	ttgggctgag	gtctagtgac	ttacccagtt	268
ctgtgtgtgg tagcactggt	ccccttgcag	tagcatgttt	cttttgatcc	cgtgccttgg	328
tgcagctgtg gggcagagac	actaagtatg	gtcctcaaag	agagtgccta	tgcctggggg	388
tcaaggaaca tttccagggc	tctttgcaca	gaagtgtcct	gccttctccc	aaggcttgtc	448
ttgtgatgga acatggaggc	ccagtggcgg	agggtagaga	gagagtgtag	ggacacatct	508
gtgtaatact cagagggtgt	tgagagcaaa	tgatgtgtgt	gtatgtgttt	gggaggggtg	5 <u>é</u> 8
tggttgctgc tgtggctgtt	gttgtttaaa	gaattgggga	ctgaaccctg	ggacatgtaa	628
gtcaaaatac tttagcaact	gagctgtatc	ccccacttgc	cataatgtga	gtcttaaatt	688
ttttcttgat ggatttagtt	tttcttctgc	catttctttt	ggggccagga	gccagtgtgt	748
atgtetgttg ccageettgt	tttatgtggt	gggcactggg	tggaacttta	ttcaaaccag	808
gggaattggg gaaatagaaa	actgttaccg	ggctgcataa	gtgaagggtg	atttgtttct	868
agatagcaag gaccttctgg	attgctttgg	gtgaaagtga	aagttgtttg	aagccatgtt	928
ccagagagag cattgcacag	tgatgtcatg	gtccatagag	aacactgcac	agtgatgtca	988
tggtccaaag ggagcactac	acagtgatgt	cacaatccgg	agagagcact	gcacagtgat	1048
gtcacagtnn nnnnnnnnac	acagtgatat	cacagtccat	agagagcact	gcacagtgat	1108
gtcacagtcc atagagagca	ctgcacagtg	atgtcacggt	ccaaagggag	cactacacag	1168
tgatgtcaca gtccatagag	agcactgcac	agtgatgtca	tggtccaaag	ggagcactac	1228
acagtgatgt cacggtccat	agagagcact	gcacagtgat	gtcacggtcc	atagggagca	1288
ctgcacagtg atgtcacggt	ccatagggag	cactgcacag	tgatgtcacg	gtccataggg	1348
agcattacac agtgatatca	cagtccatag	agagcactgc	acagtgatgt	catggtccat	1408
agggagcact gcacagtgat	gtcatagtcc	atagagaaca	ctacacagtg	atgtcacggt	1468
ccatagggag cactgcacag	tgatgtcaca	gtccatagag	agcactgcac	agtgatgtca	1528
cagtcaatca ataatactgc	acgatgattg	tcacaccaat	cccctttgcc	agagaccaca	1588
ttcaggaagg caggggagca	ggacatcggg	getetecate	cgtgtgcttg	ttcgtccctc	1648
tacccacttg gtcaccgaat	tgatgcactc	attgcgtctg	ctcctacgcc	ccaccatctc	1708
ctctcaccta cctgctcact	tgtgtattgt	gttagttacc	ttttgattct	gtgattggat	1768

ctcacaaaag caacttcagg caaaggcttg ctttggctta tgatttaaga ggagatatag 1828 tctaacacgg aaggaaggca catggcagga gcaggagacc gtgggtcaca atcatgggag 1888 atgagtgctg gtgtccatct cagtttactt tttattcagc ccgggaccct gacctgggat 1948 gatgccaagt acatttattg cgtggctctt cccatctcaa tccaatgtag ccgttccctc 2008 acagagaggg cgctcagagg cctgccttct gggtgcctct tgaccccgtt ctgctgacac 2068 ccacacatgc gctcacccag gtattttcca tttaatggct cattcgccca ttcatttgtt 2128 cagccgctca agtggtattt gcagacggct cttccttgta agggggcgtg caggaggggc 2188 agaggeteag ageaetgaet getgtggaag atacaageet eteeteteae etgeeaeage 2248 tgtgaatggg cagttgtcct tgaaggagtg agtggtggcg tcctgtgcct tagggatgga 2308 caagaaactg cagctgtgtg cagttcctca ggcccccttg tctgcctctc ctaggttgtg 2368 gtcagcgttt tgctgtggga tagttgggag aatgaagtca ggctcctgtg gttagactgg 2428 gcatgctaag ctttctggta gaccctgggg gctgaagtca ggacaggcat gcttaggtct 2488 ctgctggagg actagaggat gtgaccaggc tgatggctag agtcctctgt ctcaagctcc 2548 taagettetg egteettgae etgatgeeea geacaagatg etcagtaaae teegggaagg 2608 ggtagtcagg ctttcttctg ctgggttaga tccccattgc ttctgtcctt cagcctctcc 2668 tetgegeeta gagacagata gatagaetge aggtgtette agtteeteag etgeeettgt 2728 ctgcccctca cagtgtgttc tgaggttgtg cctggagttg tagccaggga agtttcacag 2788 tattgcatca gggtgctggg acagaggtct tttccgttgg ggctgtttgg actgacgtcc 2848 tetetgagte tattteeeat tgetgtgaga aaceteeeat ttttacatag tteataagea 2908 geacttattg ttgttgtgea ttaegettga etaetttgae tetgtagaeg attetetgte 2968 tatattcata tgttttcccc accatgccca agactcttac tgagctgctc aggctcaagg 3028 atattgcact agetetgtgg ttaggtatgt cacegaggta gggtgcagtg gggagaetgg 3088 actttgtcat ataaagggca ggagagactc cagcggcaaa atgactcatt tgatcttcaa 3148 gattaccctg gctatgacct caacagccat tgctatttct ggaatatttt cccaaggtgc 3208 aaagaacaga tggtctgtgg gactcactca cactgaaggc cagctgtgaa gggcagcatg 3268 gtgcttgccc tcatgaactg catacccagt tcctgccatc gactgtaacg aagaaggggt 3328 aacgggcaca gtggtgctta gcctgaggac agacttggga gcgtgcacag tgagagtgta 3388 gtctgtcttc taacttgaga gctagagagc taggggccct ggatgtcacc agcaagttga 3448 agagatggcc ttgtggtagg ggtgacacca tggccaaacg gcatccagat caagccgctt 3508

aaagatcagg tgaacttcaa agtgctttga gactggtgct tgttccctgc agaccccttt 3568 aggaggtgga tggcttgaca gagcactcaa acaggtcaaa gctggcagag caccatgcta 3628 aagetgggte etgeatttga acaggettae ggtetgtett aattteetat gtgaccatgt 3688 gattttctca tcatttgggt aaaactttga aacgacaaac tgtcagactt tctaacattt 3748 gtgtatgtgt ctgcacaagc acagatgtgt gtgagcatat gcatgtgtgt atatgtggat 3808 gagcctgtgt gtcctctgca tgtgcacaca aaggcacatg catatgtttg tgtatccatg 3868 ttagtttgtg tgtgtgtgtg tgtgtgtgat acacatctgt gtatggtaag tatacatgtg 3928 tgttgtgtgc atgtgtgagt cattccatgt atttgtgcat tcctgttagg tgtttatgta 3988 ccacctgtac gttgtgtgaa cgtgtacatg ctgtgggata ctagaaaggc atggagcttc 4048 tgggagetgg agetgetett teeceggeet eeettttgtt getteaaaae aetgtgttga 4108 agcatgggga ccgaaagtga aagccatgct taccggtcgt ccatctgctg aggaaacact 4168 teetgageat etgetetgte etgggeetgg ggeeecaaag gtgaaagtgt aaataggget 4228 gtgagatagg acagttagat gttcacagct tggggccagg gcacctttga ggagggaagg 4288 atcacgtgca cctaagctgt gtgaaaggag gcaagatgaa cggagtgtca ggcaactctg 4348 aggcaggtac agagagaggg agaactggga agatcgttga ggcttgagtg aaaatgttaa 4408 agetgggaga gattgagaga eettgaaaet gaagttaaae aggtaatttt gettttgttt 4468 tacttgggaa aatattccca aacagggtta acgtgtatca tgtccacgca tttccactag 4528 tgttaaagaa tggacttgat gagtagttgg ccccttggaa ttgtccctag gatccaacaa 4588 gattagcaag tatgtgaggc ccaagtagaa tatcctcttt tctggaagca aactgagcac 4648 gacttaaagc ttgatgagca gcataggaga aagaagggtc tgctgtgcca aagggtatca 4708 tccccgtgag gagggetttt ctagttaggc cagcaaactc ccagggaaca ccagtgtgtt 4768 cccagggaga tgggaacaat cgagaggcat ctaggtgtgt attcctcagt agggtttctg 4828 tgtctgagaa atggagggga cacatcacag gagtaagggc cagggtgaga gggagacaaa 4888 gagaaagget etggeattea ggaactgagg geaggaeaga eageaagtee agttgggaet 4948 ggtcagtggt ggggctcatg gtgggcgtgg ggtgtttgtt cttctgcatt attctgagta 5008 gttcctgctt agcctcacat ccaggcatta tcttggggct gacactttct ggggagcact 5068 ggacttcagg tcccactagg gagtagcctg tgtagctgta cccctcacct ccacctcaca 5128 geteatette cattgeagee tgggtgggtg ggggaegggt tecateetge etgttgeegt 5188

aacctgtgac atttcctttc cag t tgc cgg att cat tat aag gat atg tac  Cys Arg Ile His Tyr Lys Asp Met Tyr  45 50	5239
agt ttg ttg cgt tgt att gcg cca ccc gtt ggc tta ggg aag aac tgc Ser Leu Leu Arg Cys Ile Ala Pro Pro Val Gly Leu Gly Lys Asn Cys 55 60 65	5287
cct cgt agg ttg gcc tac aag gtttgtaact tcagagcctc catctgacat Pro Arg Arg Leu Ala Tyr Lys 70 75	5338
accecegee egaategeae acaetgeaea caetecetge etgaettggg aatgacaaca	5398
cctcaccttc tetttectct gtcctgecct gccctgcgtt ccagcatect ggatgetttg	5458
gttccctgtc ttatgtcccc tagctggagg caggcagcct tcatgacagt gtctgagatg	5518
gtttctggcc tggctcggcg gagtctgcca ggaccaattg gctgggcgtg ggggttaggg	5578
agggggttgg catgtggttg agacaggtcc cacctttacc aagcaaacag ccatcacacc	5638
cagcaatcct gcaggctctt atgattgggc tgatggtccg acagccagag gctgagagcc	5698
gagaggagaa tttgtacatg ggtgcaaata aacaaggagc agttgtccct gtgggtcagg	5758
agaggetete aggteeetgg gggaggggag agtgggaaca agaaagggga geeeecetge	5818
gtcgcttccg gattgatgtg ctcagctcag ggagaaaacc cctggcctgg	5878
gaagggcagg gtctgggcag atctgccctc tggagaaagg ctggactctg ctgtgtctct	5938
gagaccccaa gcccacaggc aggtgctctg gggtagaggg aagccccagc ctcttgactt	5998
gggaagatgc tacaccagcc ccaagatgct gccctgagtg ccccagccca caggggcgca	6058
tccaacctgt tggtcccaac ctgtgctgtc agaggcacta gtggcaccag gccacagctt	6118
actcagctcc tecetteceg cagteteett eccatecage ectgaettea geageagece	6178
tgccctgggc tggcctcggg cgtctctggg cttggagctg acttgccctg tcattcagtt	6238
cagtcactaa gaactttggg actccctttc tagctttcta atctgttgta gacttttctc	6298
cacattagee teeggeetta tgtggaagea aateeaatet geecattgag gtgggegaea	6358
ctgctgctgt ggttctcagc ggtagtctcc tctccacagt ctggccagtg cttcccgtgc	6418
catgaaccag ctctttctga tttggccttg tctgcagttc taaaacagta aagtcagaaa	6478
tttataaccc aaagccgccc cgatgcagcc cttcctcacc cctccatgtt cattagctca	6538
gacatcaccc ctgcccgctg ccctgcattg ggctgcctat gaggaggtga ggctggggt	6598
gggggcaagc aagggtcaag tcaagcagag ataccttcat tttacttgtt tttgatttgg	6658
ggcatttacc cgaactttgg ttaactgaat ctgttttctt gtctgcttcc tcccctgccc	6718

gctccctgct ctgcccctct gccgtgcgcc tctcctgtgg accaccccac tatgcctggc	6778
ctctggaacg ggtttccag tgg gcg cat cag tta caa tga cat gtt tga gat Trp Ala His Gln Leu Gln His Val Asp 80	6830
gct gaa aca cat gtc ccc acc tct ggg ttt ggg gaa gaa atg ccc ggc Ala Glu Thr His Val Pro Thr Ser Gly Phe Gly Glu Glu Met Pro Gly 85 90 95 100	6878
tcg agt tgc ata caa g gtagaccttg accctgcact gatcctgcat ccaacccagg Ser Ser Cys Ile Gln 105	6934
agetgtetgt gettggetgg geceaectgg eceteattea geceaeagag eacteatggg	6994
ctctcacctg cttccttagg atttgtctgg ctgtctggct gtacctggct gccttactgc	7054
tgetecetea ecacecegaa ateetttget eccateceag acagaggeat aegtetetge	7114
actaatccac taccctgtct gacttcccca ccactgacat ccttgaggag cttgggacaa	7174
tggagccaga gataccccat ccagggccct tttcccttct ttgcctgtgc tttcccaaga	7234
atgagaagac caacaaaaca atgcttttgg accgagggtg ctttcctcct agatgccctg	7294
cccctgtgct cctgtcatgg gtggtcatca agtctgcaag gtctgcctcc agggacagat	7354
ggggcctgcg ggcctctgct gctcccctgt gccgtgccct tctaccccag ctaagctcag	7414
ccttgtaggg tttgtgccca gctagtcttc aagggcttca aaacttccca gtgggaagga	7474
ggtggggcct tgacgttgga agggaaaggc tgcttggatg ccttggttaa ggctccagct	7534
tcgcattcag gagtcccagc acacgggagt cgtgaggaag gaagtgaaca tggacacctc	7594
caggeeteet teetggeeca gteatagete getgeacetg cagtgeeetè ageacteata	7654
gcccctactg ctccccacta aacagtgcca actgtgttag catctgtctg agagctccct	7714
gctggctcct ccctcttcag cccgctgtac tctgtccatc ttctgtctcc gtaccccatt	7774
ctaggttttt ttgtctgcca cttcattcag tgtagtctgt gggaaccaaa cagctctccc	7834
accactcaca gatgeetetg ggttetteet geetttatet geececaact ageatgtgtg	7894
ttgaaactgc ggaggcettc tgcatgcatg cgtgcatgtc tctacccaca ggagggagag	7954
ctctgttcct atgacaccca gtcagacaga aaggcaggca ggaagggttg cactgccctt	8014
tccagccagg cctgaggatc ctgaggttca gggtgtgacc agaccaagag aggcttcggg	8074
gtaggttgca gatggcttgg ggcaaaatgg gggagggccc aagagatggt tccagtacca	8134
tttctggaac ttatttgcca tggtcttgcc actctgcttc tctgaggttc tatctttctc	8194

tgtcctccat	gactgggggc	aaacctgtgg	gagcccctgt	gagggtgacc	tttggtgtcc	8254
tggacagatc	tcccctctt	ggaaacatct	gcttgctctt	ctctcattcc	catcctgacc	8314
cctccctgga	agaggaattt	cttggcctct	cacactgcca	tcggtgctgg	tctgcaggga	8374
cgccatgctc	cgggaagcag	gcttttcctg	gtgtgtaaca	catgtgacac	gtgaggtcct	8434
ggtcctgtga	ttggacgtgc	ccagtggggc	tttgccagca	ttcagaggac	aagagcctac	8494
ctcctgtgtt	ctgcacccca	tgcccctcta	tcacaatgaa	gatacagaac	gtgggggcaa	8554
gatectaggg	accgggctgt	gtggcccttc	cctgaagaag	ggtgcttgcc	ttagggagga	8614
agtcctatct	ctagttttct	tcctatacag	aggacattag	tattcctggg	gtgcttcatg	8674
gccgactgag	tggtntcaga	tgcacctgag	acctccctca	gatgttntga	gagccggact	8734
tccttgcagc	tgctgagtct	acccagagga	atgactcaga	cacatccctg	tttcaagatg	8794
tgttgtctct	gaaaaacaag	aaaagaaagg	gagctggcca	ctcagagata	taggagcagc	8854
ttgctctgct	ctggccctta	gaaaggtctc	ttaagactct	tcttcattca	tttacctatg	8914
atggtgtctg	tcttcaagtg	tgtctccccg	gccccctgtt	tgtggttaag	acctgagtta	8974
agcaagagca	tctctcttga	tgagtgcctc	gttgggactg	gacctcagag	aggcctcttg	9034
gcttccaccc	cttacttacc	caatgatcca	gttgcagata	gagggcaggt	aagcaaaaac	9094
cacaggacag	ggcctccaac	tgtattctaa	gctgtcccac	tetgeettee	tcagccagga	9154
ccatgccccc	ttgtgtcact	catagccagt	ggtcaggact	caggeteetg	ggcctggctt	9214
ctgctacagc	ttctacttgg	acttctgttc	ttctctagtc	aggagaatct	cagcttagat	9274
gagagtaagg	tccaggagga	gagaatgctg	gcaaaacttc	caccgcccag	ccttctggac	9334
caggcacctg	ccaggaccac	tctcgggaca	ccaagccaag	ctccagagcc	cagacactaa	9394
ctctagggtg	ggagtgaccg	taatatgctt	ggccctgatc	tgcccctcct	catccccag	9453
			cat atc caa His Ile Gln 115			9500
aca ctt tac Thr Leu Tyr	atc cac ac Tle His T 125	ct gat ggc o nr Asp Gly I	cct cat ccg Pro His Pro 130	gac ggc act Asp Gly Thi	gga gat Gly Asp 135	9548
caa gct tgc Gln Ala Cys	_		·			9562

<210> 24 <211> 334

<212> <213>	DNA Ratt	us no	orveg	icus	3											
<400> atgctga	24 aacc	tcttt	gttg	c to	gtgat	cato	g gad	caatt	tcg	aata	accto	cac c	geggg	gattct	;	60
tccatco	ctag	ggccg	gcacc	a co	ctcga	atgaa	ttc	catto	gcg	tcts	ggct	ga a	ataco	gaccca	a :	120
gctgcgt	gtt	gccgg	gatto	a tt	ataa	aggat	ato	gtaca	igtt	tgtt	gcgt	tg t	atte	gegeea	<b>a</b> :	180
cccgtt	ggct	taggg	gaaga	a ct	gccc	ctcgt	agg	gttgg	gcct	acaa	agcgo	ect g	ggtto	gaatg	3 :	240
aacatg	cca	tatco	caatg	a gg	gacat	gace	g gta	acact	tta	cato	ccaca	act ç	gatgo	geceto	2 :	300
atccgga	acgg	cacto	ggaga	it ca	agct	tgcc	c cca	ag							:	334
<210> 25 <211> 111 <212> PRT <213> Rattus norvegicus																
<400>	25															
Met Le	ı Asn	Leu	Phe 5	Val	Ala	Val	Ile	Met 10	Asp	Asn	Phe	Glu	Tyr 15	Leu		
Thr Arg	g Asp	Ser 20	Ser	Ile	Leu	Gly	Pro 25	His	His	Leu	Asp	Glu 30	Phe	Ile		
Arg Va	l Trp 35	Ala	Glu	Tyr	Asp	Pro 40	Ala	Ala	Cys	Cys	Arg 45	Ile	His	Tyr		

Lys Asp Met Tyr Ser Leu Leu Arg Cys Ile Ala Pro Pro Val Gly Leu 55

Gly Lys Asn Cys Pro Arg Arg Leu Ala Tyr Lys Arg Leu Val Arg Met 65 70 75

Asn Met Fro Ile Ser Asn Glu Asp Met Thr Val His Phe Thr Ser Thr 85 90

Leu Met Ala Leu Ile Arg Thr Ala Leu Glu Ile Lys Leu Ala Pro 100 105

<210> 26

<211> 111

<212> PRT

<213> Rattus norvegicus

Met 1	Leu	Asn	Leu	Phe 5	Val	Ala	Val	Ile	Met 10	Asp	Asn	Phe	Glu	Tyr 15	Leu	
Thr	Arg	Asp	Ser 20	Ser	Ile	Leu	Gly	Pro 25	His	His	Leu	Asp	Glu 30	Phe	Ile	
Arg	Val	Trp 35	Ala	Glu	Tyr	Asp	Pro 40	Ala	Ala	Cys	Gly	Arg 45	Ile	Ser	Tyr	
Asn	Asp 50	Met	Phe	Glu	Met	Leu 55	Lys	His	Met	Ser	Pro 60	Pro	Leu	Gly	Leu	
Gly 65	Lys	Lys	Cys	Pro	Ala 70	Arg	Val	Ala	Tyr	Lys 75	Arg	Leu	Val	Arg	Met 80	
Asn	Met	Pro	Ile	Ser 85	Asn	Glu	Asp	Met	Thr 90	Val	His	Phe	Thr	Ser 95	Thr	
Leu	Met	Ala	Leu 100	Ile	Arg	Thr	Ala	Leu 105	Glu	Ile	Lys	Leu	Ala 110	Pro		
<210 <211 <212 <213	l> 9 2> I	27 9695 DNA Ratti	us no	orveç	gicus	3										
<220 <221 <222	L> (	CDS (92)	(70	93)												
<400 cgg		27 agc g	ggcta	aggtt	a gg	gagco	ccctç	g gcg	geged	egeg	ccct	cggt	ige (	eggge	ccgcgg	60
agco	29999	gat g	gegeg	gegge	cg co	ccgg	ggagt	1		-	ege t Arg I	he (		-	-	112
											gag Glu					160
											999 Gly 35					208
											caa Gln					256

_	gaa tgo Glu Cys	Arg G	-										976
	gac aac Asp Asn		_	_		_				_	_		
_	gag ggc Glu Gly 315	Trp T	_							_		_	1072
	acg tgg Thr Trp 330												1120
tcc ttc Ser Phe 345	ttc atg	g ctc a : Leu A	ac ctg sn Leu 350	gtg Val	ctc Leu	ggt Gly	gtg Val	ctt Leu 355	tca Ser	gga Gly	gag Glu	ttt Phe	1168
_	gag cgg Glu Arg	g Glu A				_	_	_		_	_		1216
	cag cag Gln Gln	_		_	_	_				_			1264
	aag gcg Lys Ala 395	ı Glu G											1312
	aag too Lys Ser 410				_	_	_	_	-		_	_	1360
	aat gac Asn Asp												1408
	tgt gct Cys Ala	Ala G											1456
	aca gag Thr Glu												1504
	ctt ato Leu Ile 475	Arg A											1552
	tgc gtg Cys Val 490												1600

	aat Asn 505															1648
	ttc Phe															1696
	G1 y 999															1744 .
	gtg Val				_			_	_	_		_	_		_	1792
	gga Gly															1840
	att Ile 585			_		_					_			_	_	1888
	tcc Ser															1936
	ttc Phe					_		_	_	_		_	_	_		1984
	gga Gly															2032
	ttc Phe		_	_						_		_				2080
Asp	tgg Trp 665	Asn	Ala	Val	Met		His	Gly	Ile	Glu	Ser	Gln				2128
_	aaa Lys		_								_	_		_		2176
	aac Asn			_	_		-		_	_			-	_		2224
	gcc Ala															2272
gca	gcc	aat.	cag	aag	ctt	ġct	ctt	cag	aag	gcc	aaa	gaa	gta	gct	gaa	2320

Ala	Ala	Asn 730	Gln	Lys	Leu	Ala	Leu 735	Gln	Lys	Ala	Lys	Glu 740	Val	Ala	Glu			
					gct Ala											2368		
	-				cgc Arg 765					_		_	_	_		2416		
					cgt Arg											2464		
					ctg Leu			_	_	_	_					2512		
gac Asp	atg Met	aag Lys 810	aca Thr	cac His	atg Met	gac Asp	cga Arg 815	ccc Pro	cta Leu	gtg Val	gtg Val	gaa Glu 820	cct Pro	ggt Gly	cgg Arg	2560		
					ccc Pro											2608		
					gcg Ala 845											2656		
					tca Ser											2704		
agg Arg	aca Thr	gac Asp	tgc Cys 875	cca Pro	aag Lys	gcc Ala	gaa Glu	agc Ser 880	acc Thr	gag Glu	acc Thr	gly aaa	gcc Ala 885	cgg Arg	gag Glu	2752		
					cgt Arg											2800		
gac Asp	aca Thr 905	caa Gln	gtg Val	cgt Arg	tgt Cys	gag Glu 910	cgc Arg	agt Ser	aga Arg	cgt Arg	cac His 915	cac His	cgg Arg	cgc Arg	gga Gly	2848		
tcc Ser 920	ccg Pro	gag Glu	gag Glu	gcc Ala	act Thr 925	gaa Glu	cgg Arg	gaa Glu	cct Pro	cgg Arg 930	cgc Arg	cac His	cgt Arg	gcc Ala	cac His 935	2896		
cgg Arg	cac His	gca Ala	cag Gln	gac Asp 940	tca Ser	agc Ser	aag Lys	gaa Glu	ggc Gly 945	aag Lys	gag Glu	ggc Gly	act Thr	gca Ala 950	ccg Pro	2944		
					ggc Gly											2992		

955 960 965

acg ggc ccc cgt gag aca gag aac agt gag gag ccc aca cgc agg cac Thr Gly Pro Arg Glu Thr Glu Asn Ser Glu Glu Pro Thr Arg Arg His 970 975 980	3040
cgt gca aag cat aag gtg cca cca aca ctt gag ccc cca gag agg gag Arg Ala Lys His Lys Val Pro Pro Thr Leu Glu Pro Pro Glu Arg Glu 985 990 995	3088
gtt gca gag aag gag agc aac gtg gtg gaa ggg gat aag gaa act Val Ala Glu Lys Glu Ser Asn Val Val Glu Gly Asp Lys Glu Thr 1000 1005 1010	3133
cga aat cac cag ccc aag gaa cct cgc tgt gac ctg gag gcc att Arg Asn His Gln Pro Lys Glu Pro Arg Cys Asp Leu Glu Ala Ile 1015 1020 1025	3178
gcg gtt aca ggc gtg ggc tct ctg cac atg ctg ccc agc acc tgt Ala Val Thr Gly Val Gly Ser Leu His Met Leu Pro Ser Thr Cys 1030 1035 1040	3223
ctc cag aaa gtg gac gaa cag cca gag gat gca gac aac cag cgt Leu Gln Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg 1045 1050 1055	3268
aat gtc acc cgg atg ggc agt cag ccc tca gac ccc agc acc act Asn Val Thr Arg Met Gly Ser Gln Pro Ser Asp Pro Ser Thr Thr 1060 1065 1070	3313
gtg cat gtc cca gtg aca ctg aca ggc cct ccc ggg gag gcc act Val His Val Pro Val Thr Leu Thr Gly Pro Pro Gly Glu Ala Thr 1075 1080 1085	3358
gta gtt ccc agt gct aac acg gac ctg gaa ggc caa gcg gag ggc Val Val Pro Ser Ala Asn Thr Asp Leu Glu Gly Gln Ala Glu Gly 1090 1095 1100	3403
aag aag gag gca gag gct gac gat gtg ctg aga aga ggc ccc agg Lys Lys Glu Ala Glu Ala Asp Asp Val Leu Arg Arg Gly Pro Arg 1105 1110 1115	3448
ccc atc gtt ccc tac agt tcc atg ttc tgc ctc agc ccc acc aac Pro Ile Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr Asn 1120 1125 1130	3493
cta ctc cgt cgc ttc tgc cat tac att gtg acc atg cgg tac ttt Leu Leu Arg Arg Phe Cys His Tyr Ile Val Thr Met Arg Tyr Phe 1135 1140 1145	3538
gag atg gtg att ctt gtg gtc atc gcc ttg agc agc att gcc ctg Glu Met Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile Ala Leu 1150 1155 1160	3583
gct gct gag gat ccc gtg cgg acc gac tca ttc cgg aac aat gct Ala Ala Glu Asp Pro Val Arg Thr Asp Ser Phe Arg Asn Asn Ala 1165 1170 1175	3628

	Lys				tac Tyr 1185										3673
	Val				ata Ile 1200										3718
	Tyr				ctg Leu 1215										3763
	Gly				gca Ala 1230										3808
	Ile				aag Lys 1245		-	-	-	_	_	_	_		3853
	Leu				aag Lys 1260										3898
_	Cys		_		tct Ser 1275	_	_		-	_			_		3943
-	Tyr	_			atg Met 1290				_	gtc Val 1295		_	gtc Val		3988
ctc Leu 1300	Phe				ttc Phe 1305										4033
	Glu				agg Arg 1320		-		_	_			_	_	4078
	Val	_		-	cca Pro 1335		_		_			_			4123
	Asp				tgg Trp 1350										4168
	gga Gly				ccc Pro 1365					cac His 1370					4213

tcc Ser 1390					gtc Val 1395										4303
gtc Val 1405					gcc Ala 1410	_						cag Gln		_	4348
999 Gly 1420					tct Ser 1425										4393
					gcc Ala 1440										4438
atg Met 1450					cag Gln 1455										4483
gtg Val 1465					ttt Phe 1470					atg Met 1475	_	_		_	4528
					ctg Leu 1485										4573
gag Glu 1495					ctg Leu 1500										4618
atg Met 1510					tgc Cys 1515										4663
ttg Leu 1525					gat Asp 1530										4708
gtt Val 1540					act Thr 1545										4753
acg Thr 1555					aac Asn 1560									gca Ala	4798
					ctg Leu 1575										4843
ttg Leu 1585					gtc Val 1590										4888
tgc	ctc	ctc	att	gcc	atg	ctg	ttc	ttc	atc	tac	gcc	atc	atc	ggc	4933

Cys 1600	Leu	Leu	Ile	Ala	Met 1605	Leu	Phe	Phe	Ile	Tyr 1610	Ala	Ile	Ile	Gly	
atg Met 1615	Gln				aac Asn 1620										4978
aac Asn 1630	Arg				ttc Phe 1635										5023
ttg Leu 1645	Phe				act Thr 1650			_			_		_	_	5068
tct Ser 1660	Cys	_			cgg Arg 1665	-	_	_			_		_	_	5113
_	Cys		_	_	ttt Phe 1680	_					_				5158
ttc Phe 1690					ctg Leu 1695	_	_				_	_			5203
atg Met 1705	Asp				tac Tyr 1710										5248
ccg Pro 1720					gaa Glu 1725										5293
cca Pro 1735					cgc Arg 1740										5338
ctg Leu 1750					cca Pro 1755										5383
gct Ala 1765					aag Lys 1770										5428
tcc Ser 1780					acg Thr 1785							ctg Leu	_	_	5473
ctc Leu 1795				-	ctg Leu 1800			_				gċg Ala			5518
aaa Lys					gat Asp										5563

1810	1815	1820
		g gac tta ctg gtg cca 5608 u Asp Leu Leu Val Pro 1835
		g aag gtc tat gcg gct 5653 y Lys Val Tyr Ala Ala 1850
	_	c aaa acc acc aga gat 5698 n Lys Thr Thr Arg Asp 1865
		c cag atg ggt cct gtt 5743 r Gln Met Gly Pro Val 1880
		g gag cag aca cag ccc 5788 u Glu Gln Thr Gln Pro 1895
		t cga caa aag agt gca 5833 u Arg Gln Lys Ser Ala 1910
<del>-</del>		a acc cag gaa agt ggc 5878 n Thr Gln Glu Ser Gly 1925
	tcc tgg ggc acg cag Ser Trp Gly Thr Glr 1935	g agg acc cag gac gta 5923 n Arg Thr Gln Asp Val 1940
		t ggc cat tct gca gag 5968 g Gly His Ser Ala Glu 1955
		t gta gat gtc cag atg 6013 a Val Asp Val Gln Met 1970
		g gag ccc cag cct ggc 6058 y Glu Pro Gln Pro Gly 1985
		g cca cgc ctg gcg gca 6103 t Pro Arg Leu Ala Ala 2000
		c atg aag cgc tcc atc 6148 o Met Lys Arg Ser Ile 2015
2 2	3 3 333	t cag ctt tgc aac aca 6193 r Gln Leu Cys Asn Thr 2030

_	Leu	_					_	_		tcc Ser 2045	His			cac His	6238
										cag Gln 2060					6283
	Gly									ggt Gly 2075	Ala				6328
-	Ala				_				_	ggg Gly 2090				_	6373
cgg Arg 2095	Arg	gag Glu	cgt Arg	aag Lys	caa Gln 2100	gag Glu	cga Arg	ggc Gly	cgg Arg	tcc Ser 2105	Gln	gag Glu	cgg Arg	agg Arg	6418
_	Pro							_	_	cgc Arg 2120				-	6463
_	Arg			-						cct Pro 2135	Lys			ctc Leu	6508
	-				_				_	cta Leu 2150					6553
cac His 2155										agc Ser 2165	Pro			tca Ser	6598
aca Thr 2170										999 Gly 2180					6643
	Gln	Thr	Pro	Leu	Thr	Pro	Arg	Pro	Ser	atc Ile 2195	Thr		_		6688
gcc Ala 2200	aat Asn	tcc Ser	tcg Ser	cct Pro	gtc Val 2205	cac His	ttt Phe	gct Ala	gag Glu	ggt Gly 2210	cag Gln	agt Ser	ggc Gly	ctt Leu	6733
	Ala									ggc Gly 2225	Leu				6778
										cag Gln 2240					6823

ggc tcc cgc att ggc tct gac cct tac cta ggg cag cgt ctg gac Gly Ser Arg Ile Gly Ser Asp Pro Tyr Leu Gly Gln Arg Leu Asp 2245 2250 2255	6868
agt gag gcc tct gcc cac aac ctg cct gag gat aca ctc acc ttt Ser Glu Ala Ser Ala His Asn Leu Pro Glu Asp Thr Leu Thr Phe 2260 2265 2270	6913
gaa gag gcc gtg gcc acc aac tct ggc cgc tcc tcc agg act tcc Glu Glu Ala Val Ala Thr Asn Ser Gly Arg Ser Ser Arg Thr Ser 2275 2280 2285	6958
tat gtg tcc tcc ctc act tcc caa tcc cac cct ctc cgc cgt gta Tyr Val Ser Ser Leu Thr Ser Gln Ser His Pro Leu Arg Arg Val 2290 2295 2300	7003
ccc aat ggc tac cac tgc act ttg gga ctc agc acc ggc gtc cgg Pro Asn Gly Tyr His Cys Thr Leu Gly Leu Ser Thr Gly Val Arg 2305 2310 2315	7048
gcg cgg cac agc tac cac cac cca gac cag gat cac tgg tgc tag Ala Arg His Ser Tyr His His Pro Asp Gln Asp His Trp Cys 2320 2325 2330	7093
ctgcaccacg accacccatg caccagctcg tgggtgcggg ttccagttga tgagtttat	7153
catcegetet gggttgtgeg gteacageee tgggaggagg gteeteacat egeggeetet	7213
gtggtggagg ttcctgcttc tctccctccc tcccttttac actggacaga ctaataaagc	7273
cetttettag agggatatgg teetetetat eeteetgtgt aetgeettee tgggtteeat	7333
gccagatgtt ggatcctaag cagaggtagc tgagttgaga tagacccagc aaatccaaat	7393
cctatgtcat ggcctccagc ttccagggtg ggtacttggg actttcttag gaggtctgag	7453
cctcatggag attgtggttt gtccaaatgt gtggcatggg ggatagggta ccctcaaagg	7513
caaggaaagg agcccaactg tgtggcctgg cagcacctgc cagcatcact actctcatgt	7573
ctattgtggt cttggagtca aacagcacat gtatatagag atatgctcaa gggcctgcct	7633
ttcacctaca ttgtcaccat aatagggacc aaatctagag gatgtccttg ctgttgattc	7693
tggttttcag tcacaacact ttcacttttt gtcatttcta tatagttgat ctagaaaaac	7753
agaaatcaaa acagggaaga aaatgttcgt gtaacttaaa aaagaaatca acgtgtagga	7813
aggtetecat titgeatigt tietgigaet igtatgeaat giteetgiat giattetaee	7873
cttcccggga agtccccaat gaccctggtt cctctgctca accaagtgcc tgatctctgg	7933
ctctgagcat cgtggctgag gtgcggcctc aggaagcatc ggggagctgc tcagagcagc	7993
actaggactt gtgtcttagg gacactgacc gtgtccagca gcatgtcaga gaagcagctg	8053
tagtgcccat gttcctccct gagtgatggg ttctgaagaa gccagagcag cacaatgtgt	8113

gettgegtga ggeaetttee geettttaaa atetgattet eagggatggg atgeetgeea 8173 agtagggtgt gatctctgtt gtgttttaaa aaacaacaac aacaaacaaa caaaacctag 8233 tattcactga atgctgaaga gagcaaaatg caagcaaaga agggactggg gttagaggga 8293 gaagcccgca ctggcagcat aataagaaac tggcagggag gggatggtcc tggaacaggc 8353 caggtgccta gagctgagtc cagcccctgg cccggaactg gggacacagc actcaaataa 8413 aacctcatgg ctacttggtg aaaggcaaac ccatgctcag gaaggtgttc agtgtgcaga 8473 gatggctgtg aggccatgag agaaaggttt cacataggca ggcagtcctt ggtgttgttct 8533 ctgtgttttg aaacgtctga tgacttcttg gtggactgtt ggtttctacc ccatgtttct 8593 cacagaagct gtgtatatgt gtgattgcgc gtgtgattgc atgtgtgtgg tagtgtgcgt 8653 gcgtgagcat gcatgagtca taggaaatgt gtgtgtgtgt gtgtgtgtgt gtgtaggtgt 8713 gtgtacgtgt gttcagcaag tggcttttgt caaccatagg gctatgcaac aaaagacaca 8773 ttactagaaa caaaacacaa gaccaccact cggtctaggg tttcagcatg attgtgacca 8833 aaccttttat agaatttcct tatatgaagg cacaataccc tgaaacttta aagataacag 8893 agtattttat tccagtaggg taagattaaa caggaccctg gactgcatgt gactgcactc 8953 atgtacaaca gaggaggatg tgcattttga tactgttctg tctctgtccc agccccagcc 9013 cttttctctt gagtgttgaa tgtatacatt ctgtgtggaa ctacagctgc tccagacagt 9073 cctgggttgg gaatcatctt tatcccacat taacatagct ggcttttctt ccaagcactg 9133 gtacacagga aaggagacat gatgtettge tteetgaett tgggtttgtt tetgtaetgt 9193 ctcttctcaa gatgttgtct gttccccctg aaatttcata gtgagttgcc aaatttgaaa 9253 tgcaacaacc agetgtetge atetggaacc tgtcaagcag tgetgtagtt tgaaaaagtt 9313 atgtgtgcat gtaaaatata cacatatata tatatacatt atacaagtat gtgcatgaaa 9373 tgtatatett cataettttt gataeaatgt atteatttgt taatttttaa ttatatttga 9433 tataaattga aggtttgttg caaaaattta tatttaacag tgttgagaga gagagaaaga 9493 gcgagagagg gagagagag gaaagatcca atcatgcaac agaaatggga ctactttaaa 9553 aatcagteet ttgaetagtt tgetgeeetg aataatattt acaaaccaaa etttggatte 9613 tgctcttgtt tctacaatga ctttttgtat aaagcaaagt ccttggatta ataaaacaac 9673 caaaaatcaa attaaaccat ta 9695

<211> 2333

<212> PRT

<213> Rattus norvegicus

<400> 28

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Thr Gly
1 5 10 15

Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Pro
20 25 30

Gly Gln Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50 55 60

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140

Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser 145 150 155 160

Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr
165 170 175

Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 . 185 190

Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205 Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 225 220

Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile 225 230 235 240

Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 255

Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala 260 265 270

Pro Ala Arg Leu Cys Asp Ser Asp Thr Glu Cys Arg Glu Tyr Trp Pro 275 280 285

Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 290 295 300

Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 305 310 315 320

Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 325 330 335

Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 350

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365

Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 375 380

Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400

Ala Glu Glu Asp Lys Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu 405 410 415

Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu 420 425 430

- Glu Gly Glu Asp Arg Phe Val Asp Leu Cys Ala Ala Gly Ser Pro Phe 435 440 445
- Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe 450 455 460
- Arg Arg Lys Glu Lys Met Phe Arg Phe Leu Ile Arg Arg Met Val Lys 465 470 475 480
- Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr 485 490 495
- Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Gln Arg Leu Thr Thr 500 505 510
- Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 525
- Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser 530 540
- Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe Glu 545 550 560
- Val Val Trp Ala Ala Ile Lys Pro Gly Thr Ser Phe Gly Ile Ser Val 565 570 575
- Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp 580 585 590
- Asn Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 595 600 605
- Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala 610 620
- Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 635 640
- Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val 645 650 655
- Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly

560	665	670

- Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr 675 680 685
- Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe 690 695 700
- Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys
  705 710 715 720
- Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln 725 730 735
- Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750
- Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp
  755 760 765
- Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys 770 780
- Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Tyr Ala 785 790 795 800
- Ser Thr Arg His Val Arg Pro Asp Met Lys Thr His Met Asp Arg Pro 805 810 815
- Leu Val Val Glu Pro Gly Arg Asp Gly Leu Arg Gly Pro Ala Gly Asn 820 825 830
- Lys Ser Lys Pro Glu Gly Thr Glu Ala Thr Glu Gly Ala Asp Pro Pro 835 840 845
- Arg Arg His His Arg His Arg Asp Arg Asp Lys Thr Ser Ala Ser Thr 850 855 860
- Pro Ala Gly Glu Gln Asp Arg Thr Asp Cys Pro Lys Ala Glu Ser 865 870 875 888
- Thr Glu Thr Gly Ala Arg Glu Glu Arg Ala Arg Pro Arg Arg Ser His 885 890 895

- Ser Lys Glu Ala Pro Gly Ala Asp Thr Gln Val Arg Cys Glu Arg Ser 900 905 910
- Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Thr Glu Arg Glu 915 920 925
- Pro Arg Arg His Arg Ala His Arg His Ala Gln Asp Ser Ser Lys Glu 930 935 940
- Gly Lys Glu Gly Thr Ala Pro Val Leu Val Pro Lys Gly Glu Arg Arg 945 950 955 960
- Ala Arg His Arg Gly Pro Arg Thr Gly Pro Arg Glu Thr Glu Asn Ser 965 970 975
- Glu Glu Pro Thr Arg Arg His Arg Ala Lys His Lys Val Pro Pro Thr 980 985 990
- Leu Glu Pro Pro Glu Arg Glu Val Ala Glu Lys Glu Ser Asn Val Val 995 1000 1005
- Glu Gly Asp Lys Glu Thr Arg Asn His Gln Pro Lys Glu Pro Arg 1010 1015 1020
- Cys Asp Leu Glu Ala Ile Ala Val Thr Gly Val Gly Ser Leu His 1025 1030 1035
- Met Leu Pro Ser Thr Cys Leu Gln Lys Val Asp Glu Gln Pro Glu 1040 1045 1050
- Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly Ser Gln Pro 1055 1060 1065
- Ser Asp Pro Ser Thr Thr Val His Val Pro Val Thr Leu Thr Gly 1070 1075 1080
- Pro Pro Gly Glu Ala Thr Val Val Pro Ser Ala Asn Thr Asp Leu 1085 1090 1095
- Glu Gly Gln Ala Glu Gly Lys Lys Glu Ala Glu Ala Asp Asp Val 1100 1105 1110

- Leu Arg Arg Gly Pro Arg Pro Ile Val Pro Tyr Ser Ser Met Phe 1115 1120 1125
- Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys His Tyr Ile 1130 1135 1140
- Val Thr Met Arg Tyr Phe Glu Met Val Ile Leu Val Val Ile Ala 1145 1150 1155
- Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val Arg Thr Asp 1160 1165 1170
- Ser Phe Arg Asn Asn Ala Leu Lys Tyr Met Asp Tyr Ile Phe Thr 1175 1180 1185
- Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile Asp Leu Gly 1190 1195 1200
- Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu Trp Asn Ile 1205 1210 1215
- Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala Phe Ala Phe 1220 1230
- Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys Ser Leu Arg 1235 1240 1245
- Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys Arg Leu Pro 1250 1255 1260
- Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser Leu Lys Asn 1265 1270 1275
- Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met Phe Ile Phe 1280 1285 1290
- Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe Phe Tyr Cys 1295 1300 1305
- Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg Gly Gln Tyr 1310 1315 1320

- Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro Arg Gln Trp 1325 1330 1335
- Lys Lys Tyr Asp Phe His Tyr Asp Asn Val Leu Trp Ala Leu Leu 1340 1345 1350
- Thr Leu Phe Thr Val Ser Thr Gly Glu Gly Trp Pro Met Val Leu 1355 1360 1365
- Lys His Ser Val Asp Ala Thr Tyr Glu Glu Gln Gly Pro Ser Pro 1370 1375 1380
- Gly Phe Arg Met Glu Leu Ser Ile Phe Tyr Val Val Tyr Phe Val 1385 1390 1395
- Val Phe Pro Phe Phe Val Asn Ile Phe Val Ala Leu Ile Ile 1400 1405 1410
- Ile Thr Phe Gln Glu Gln Gly Asp Lys Val Met Ser Glu Cys Ser 1415 1420 1425
- Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp Phe Ala Ile Ser Ala 1430 1440
- Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn Lys Gln Ser Phe Gln 1445 1450 1455
- Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro Phe Glu Tyr Phe 1460 1465 1470
- Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu Met Met Lys 1475 1480 1485
- Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu Lys Cys Leu 1490 1495 1500
- Asn Ile Val Phe Thr Ser Met Phe Ser Leu Glu Cys Ile Leu Lys 1505 1510 1515
- Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp Ala Trp Asn 1520 1530
- Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr Asp Ile Leu

1535 1540 1545

Val	Thr 1550	Glu	Ile	Ala	Glu	Thr 1555	Asn	Asn	Phe	Ile	Asn 1560	Leu	Ser	Phe
Leu	Arg 1565		Phe	Arg	Ala	Ala 1570	Arg	Leu	Ile	Lys	Leu 1575	Leu	Arg	Gln
Gly	Tyr 1580	Thr	Ile	Arg	Ile	Leu 1585	Leu	Trp	Thr	Phe	Val 1590	Gln	Ser	Phe
Lys	Ala 1595	Leu	Pro	Tyr	Val	Cys 1600	Leu	Leu	Ile	Ala	Met 1605	Leu	Phe	Phe
Ile	Tyr 1610	Ala	Ile	Ile	Gly	Met 1615	Gln	Val	Phe	Gly	Asn 1620	Ile	Ala	Leu
Asp	Asp 1625	Gly	Thr	Ser	Ile	Asn 1630	Arg	His	Asn	Asn	Phe 1635	Arg	Thr	Phe
Leu	Gln 1640	Ala	Leu	Met	Leu	Leu 1645	Phe	Arg	Ser	Ala	Thr 1650	Gly	Glu	Ala
Trp	His 1655	Glu	Ile	Met	Leu	Ser 1660	Cys	Leu	Gly	Asn	Arg 1665	Ala	Cys	Asp
Pro	His 1670	Ala	Asn	Ala	Ser	Glu 1675	Cys	Gly	Ser	Asp	Phe 1680	Ala	Tyr	Phe
Tyr	Phe 1685	Val	Ser	Phe	Ile	Phe 1690	Leu'	Cys	Ser	Phe	Leu 1695	Met	Leu	Asn
Leu	Phe 1700	Val	Ala	Val	Ile	Met 1705	Asp	Asn	Phe	Glu	Tyr 1710	Leu	Thr	Arg
Asp	Ser 1715	Ser	Ile	Leu	Gly	Pro 1720	His	His	Leu	Asp	Glu 1725	Phe	Ile	Arg
Val	Trp 1730	Ala	Glu	Tyr	Asp	Pro 1735	Ala	Ala	Cys	Gly	Arg 1740	Ile	Ser	Tyr
Asn	Asp 1745	Met	Phe	Glu	Met	Leu 1750	Lys	His	Met	Ser	Pro 1755	Pro	Leu	Gly

- Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr Lys Arg Leu Val 1760 1765 1770
- Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr Val His Phe 1775 1780 1785
- Thr Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu Glu Ile Lys 1790 1795 1800
- Leu Ala Pro Ala Gly Thr Lys Gln His Gln Cys Asp Ala Glu Leu 1805 1810 1815
- Arg Lys Glu Ile Ser Ser Val Trp Ala Asn Leu Pro Gln Lys Thr 1820 1825 1830
- Leu Asp Leu Leu Val Pro Pro His Lys Pro Asp Glu Met Thr Val 1835 1840 1845
- Gly Lys Val Tyr Ala Ala Leu Met Ile Phe Asp Phe Tyr Lys Gln 1850 1860
- Asn Lys Thr Thr Arg Asp Gln Thr His Gln Ala Pro Gly Gly Leu 1865 1870 1875
- Ser Gln Met Gly Pro Val Ser Leu Phe His Pro Leu Lys Ala Thr 1880 1885 1890
- Leu Glu Gln Thr Gln Pro Ala Val Leu Arg Gly Ala Arg Val Phe 1895 1900 1905
- Leu Arg Gln Lys Ser Ala Thr Ser Leu Ser Asn Gly Gly Ala Ile 1910 1915 1920
- Gln Thr Gln Glu Ser Gly Ile Lys Glu Ser Leu Ser Trp Gly Thr 1925 1930 1935
- Gln Arg Thr Gln Asp Val Leu Tyr Glu Ala Arg Ala Pro Leu Glu 1940 1945 1950
- Arg Gly His Ser Ala Glu Ile Pro Val Gly Gln Pro Gly Ala Leu 1955 1960 1965

- Ala Val Asp Val Gln Met Gln Asn Met Thr Leu Arg Gly Pro Asp 1970 1975 1980
- Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg Ala Ala Ser 1985 1990 1995
- Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Ala Pro Asn Ala Ser 2000 2005 2010
- Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Pro Arg Pro His Gly 2015 2020 2025
- Thr Gln Leu Cys Asn Thr Val Leu Asp Arg Pro Pro Pro Ser Gln 2030 2035 2040
- Val Ser His His His His Arg Cys His Arg Arg Arg Asp Lys 2045 2050 2055
- Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu Ser Val Asp Thr 2060 2065 2070
- Glu Gly Ala Pro Ser Thr Ala Ala Gly Ser Gly Leu Pro His Gly 2075 2080 2085
- Glu Gly Ser Thr Gly Cys Arg Arg Glu Arg Lys Gln Glu Arg Gly 2090 2095 2100
- Arg Ser Gln Glu Arg Arg Gln Pro Ser Ser Ser Ser Glu Lys 2105 2110 2115
- Gln Arg Phe Tyr Ser Cys Asp Arg Phe Gly Ser Arg Glu Pro Pro 2120 2125 2130
- Gln Pro Lys Pro Ser Leu Ser Ser His Pro Ile Ser Pro Thr Ala 2135 2140 2145
- Ala Leu Glu Pro Gly Pro His Pro Gln Gly Ser Gly Ser Val Asn 2150 2155 2160
- Gly Ser Pro Leu Met Ser Thr Ser Gly Ala Ser Thr Pro Gly Arg 2165 2170 2175

- Gly Gly Arg Arg Gln Leu Pro Gln Thr Pro Leu Thr Pro Arg Pro 2180 2185 2190
- Ser Ile Thr Tyr Lys Thr Ala Asn Ser Ser Pro Val His Phe Ala 2195 2200 2205
- Glu Gly Gln Ser Gly Leu Pro Ala Phe Ser Pro Gly Arg Leu Ser 2210 2215 2220
- Arg Gly Leu Ser Glu His Asn Ala Leu Leu Gln Lys Glu Pro Leu 2225 2230 2235
- Ser Gln Pro Leu Ala Ser Gly Ser Arg Ile Gly Ser Asp Pro Tyr 2240 2245 2250
- Leu Gly Gln Arg Leu Asp Ser Glu Ala Ser Ala His Asn Leu Pro 2255 2260 2265
- Glu Asp Thr Leu Thr Phe Glu Glu Ala Val Ala Thr Asn Ser Gly 2270 2275 2280
- Arg Ser Ser Arg Thr Ser Tyr Val Ser Ser Leu Thr Ser Gln Ser 2285 2290 2295
- His Pro Leu Arg Arg Val Pro Asn Gly Tyr His Cys Thr Leu Gly 2300 2310
- Leu Ser Thr Gly Val Arg Ala Arg His Ser Tyr His His Pro Asp 2315 2320 2325
- Gln Asp His Trp Cys 2330
- <210> 29
- <211> 7011
- <212> DNA
- <213> Rattus norvegicus
- <220>
- <221> CDS
- <222> (1)..(7011)
- <400> 29
- atg gtc cgc ttc ggg gac gag cta ggc ggc cgc tat ggg ggc acc ggc Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Thr Gly

							ggc										96
	_				_	_	ggc Gly 40	_		_	_		_	_		Ξ	144
			-	-			atg Met	_	_						_	Ξ	192
_	_		_			-	aac Asn	_	_			-		_		2	240
		_	_	_			gct Ala	_	-			_		_		2	288
	-		_		_	-	acc Thr					_		_	_	;	336
_	_		_				gat Asp 120		_	_			_		_	:	384
_	_	_	_	_	-		tac Tyr						-			4	432
							ctg Leu									4	480
							gtc Val										528
							act Thr									į	576
-		_					ctg Leu 200	_	_						_	e	624
							atc Ile									6	672
							ttc Phe									•	720

ggc ctc Gly Leu													768	
agc aca Ser Thr	a Glu												816	
cct gct Pro Ala													864	
gga ccc Gly Pro 290		Ile '											912	
ttg acc Leu Thr 305	e Gln												960	
tac aat Tyr Asn	_		-			_				_			1008	
atc ccc Ile Pro	e Ile												1056	
ggt gtg Gly Val													1104	
cgc cgt Arg Arg 370		Lys :											1152	
ctg aat Leu Asn 385	 r Leu	_				_			_	_	_	_	1200	
gca gag Ala Glu	 _		_	_		_			_	_	_		1248 .	
ttg aag Leu Lys	 a Ala		_	_	_	_		_				_	1296	•
gaa gag Glu Glu	 			_	_		_	_	_				1344	
ttt gct Phe Ala 450	 _	Leu :	_	_		_			_		_		1392	

			_		_	atg Met						_	-	_		1440
aaa Lys	gca Ala	cag Gln	agc Ser	ttc Phe 485	tac Tyr	tgg Trp	gtg Val	gta Val	ctg Leu 490	tgc Cys	gtg Val	gtg Val	gcc Ala	ctg Leu 495	aac Asn	1488
						gta Val				-		_				1536
						gag Glu										1584
						tac Tyr 535										1632
						gac Asp										1680
						atc Ile	_								_	1728
						ctg Leu						-		_		1776
			_			ctg Leu	-	_						_	_	1824
						ttc Phe 615										1872
gct Ala 625	ctg Leu	ttg Leu	Gly	Met	cag Gln 630	ctg Leu	Phe	gjå aaa	Gly	Gln	Phe	aac Asn	ttt Phe	caa Gln	gat Asp 640	1920
						ttt Phe										1968
						gga Gly										2016
						gga Gly	_	_			_					2064
tac	ttc	atc	gtc	ctg	aca	ctg	ttt	gga	aac	tac	acc	ctg	ttg	aac	gtt	2112

· ተ	vr i	Phe	Ile	Val	Leu	Thr	Leu	Phe	Glv	Asn	Tvr	Thr	Leu	Leu	Asn	Val		
1	_	690					695		1		-1-	700		~		, <del></del>		
Ph										gcc Ala							2160	
										gcc Ala 730							2208	
										agc Ser							2256	
				_	_			_		tcg Ser	_	_		_		_	2304	
_	rp (		_		_	_	_			ctc Leu	_		_	_	_	_	2352	
CΣ			_	_		_		_	_	ccg Pro			_	_	_		2400	
·-		_	_	_					_	atg Met 810				_	_	_	2448	
										ggc Gly							2496	
		Lys		_				_		gcc Ala		_			_		2544	
	ro i									agg Arg							2592	
	nr :									aca Thr							2640	
										cgt Arg 890							2688	
										aca Thr							2736	
										ccg Pro							2784	

915 920 925

gaa cct cgg cgc cac cgt gcc cac cgg cac gca cag gac tca agc as Glu Pro Arg Arg His Arg Ala His Arg His Ala Gln Asp Ser Ser Ly 930 935 940	
gaa ggc aag ggc act gca ccg gtg ctt gta ccc aag ggc gag cc Glu Gly Lys Glu Gly Thr Ala Pro Val Leu Val Pro Lys Gly Glu Ar 945 950 955 96	
cgc gca aga cat cga ggc ccg cgt acg ggc ccc cgt gag aca gag ac Arg Ala Arg His Arg Gly Pro Arg Thr Gly Pro Arg Glu Thr Glu As 965 970 975	
agt gag gag ccc aca cgc agg cac cgt gca aag cat aag gtg cca cc Ser Glu Glu Pro Thr Arg Arg His Arg Ala Lys His Lys Val Pro Pro 980 985 990	
aca ctt gag ccc cca gag agg gag gtt gca gag aag gag agc aac Thr Leu Glu Pro Pro Glu Arg Glu Val Ala Glu Lys Glu Ser Asn 995 1000 1005	
gtg gaa ggg gat aag gaa act cga aat cac cag ccc aag gaa cct Val Glu Gly Asp Lys Glu Thr Arg Asn His Gln Pro Lys Glu Pro 1010 1015 1020	
cgc tgt gac ctg gag gcc att gcg gtt aca ggc gtg ggc tct ctg Arg Cys Asp Leu Glu Ala Ile Ala Val Thr Gly Val Gly Ser Leu 1025 1030 1035	-
cac atg ctg ccc agc acc tgt ctc cag aaa gtg gac gaa cag cca His Met Leu Pro Ser Thr Cys Leu Gln Lys Val Asp Glu Gln Pro 1040 1045 1050	
gag gat gca gac aac cag cgt aat gtc acc cgg atg ggc agt cag Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly Ser Glr 1055 1060 1065	
ccc tca gac ccc agc acc act gtg cat gtc cca gtg aca ctg acc Pro Ser Asp Pro Ser Thr Thr Val His Val Pro Val Thr Leu Thr 1070 1075 1080	
ggc cct ccc ggg gag gcc act gta gtt ccc agt gct aac acg gad Gly Pro Pro Gly Glu Ala Thr Val Val Pro Ser Ala Asn Thr Asp 1085 1090 1095	
ctg gaa ggc caa gcg gag ggc aag aag gag gca gag gct gac gat Leu Glu Gly Gln Ala Glu Gly Lys Lys Glu Ala Glu Ala Asp Asp 1100 1105 1110	
gtg ctg aga aga ggc ccc agg ccc atc gtt ccc tac agt tcc atc Val Leu Arg Arg Gly Pro Arg Pro Ile Val Pro Tyr Ser Ser Met 1115 1120 1125	-
ttc tgc ctc agc ccc acc aac cta ctc cgt cgc ttc tgc cat tac Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys His Tyr 1130 1135 1140	

									- 1							
·												ctt Leu 1155				3474
	_	_	_	_		_	_	_	_		_	ccc Pro 1170				3519
			Phe									gac Asp 1185				3564
												atg Met 1200				3609
												gac Asp 1215				3654
												gtg Val 1230				3699
												gac Asp 1245				3744
												ccc Pro 1260				3789
			Arg									gac Asp 1275				3834
												gtc Val 1290				3879
	ttc Phe	atg Met 1295	ttt Phe	ata Ile	ttt Phe	Ala	gtc Val 1300	Ile	gcc Ala	gtc Val	caa Gln	ctc Leu 1305	ttc Phe	aaa Lys	Gly ggg	3924
												ctg Leu 1320				3969
	_			_		-	-			_	_	gag Glu 1335	_	_	_	4014
												tat Tyr 1350				4059

						ctg Leu 1360									4104
						cac His 1375									4149
						ttt Phe 1390									4194
						ttc Phe 1405									4239
						acc Thr 1420									4284
						gaa Glu 1435									4329
	gcc Ala 1445					ccc Pro 1450							cag Gln		4374
						aag Lys 1465	Thr								4419
						atg Met 1480									4464
						tac Tyr 1495	Asp								4509
Met	Leu	Lys	Cys	Leu	Asn	atc Ile 1510	Val	Phe	Thr	Ser	Met	Phe		_	4554
						atc Ile 1525									4599
														agt Ser	
Ile	act Thr 1550	gat Asp	att Ile	tta Leu	gta Val	acg Thr 1555	gag Glu	att Ile	gcg Ala	aac Asn	aac Asn 1560	ttc Phe	atc Ile	aac Asn	4689
ttg	agc	ttc	ctt	cgc	ctc	ttc	cgg	gca	gca	cgg	ctg	atc	aag	ctc	4734

Leu	Ser 1565	Phe	Leu	Arg	Leu	Phe 1570	Arg	Ala	Ala	Arg	Leu 1575	Ile	Lys	Leu		
_	_	_					Arg		_		tgg Trp 1590			_	4779	
cag Gln	tcc Ser 1595	ttt Phe	aag Lys	gcg Ala	ctg Leu	ccc Pro 1600	tac Tyr	gtg Val	tgc Cys	ctc Leu	ctc Leu 1605	att Ile	gcc Ala	atg Met	4824	
		Phe					Ile				gtt Val 1620				4869	
	_		_	_			_			_	cac His 1635				4914	
			_		_		_	_	_		agg Arg 1650	_	_		4959	
											ctg Leu 1665				5004	
_					_		_	_	_	_	999 Gly 1680	_	_		5049	
-					_						tgt Cys 1695			_	5094	
											aat Asn 1710				5139	
											cac His 1725				5184	
ttc Phe	att Ile 1730	cgc Arg	gtc Val	tgg Trp	gct Ala	gaa Glu 1735	tac Tyr	gac Asp	cca Pro	gct Ala	gcg Ala 1740	tgt Cys	Gly aaa	cgc Arg	5229	
											cac His 1755				5274	
	_		_		_		_	_	-	_	gtt Val 1770	_		~	5319	
											gag Glu				5364	

gta cac t Val His I 1790		tcc aca Ser Thr	_	_	_				_	_	_	5409
gag atc a Glu Ile I 1805	-	-					_			_	_	5454
gct gag o Ala Glu I 1820		aag gag Lys Glu										5499
cag aag a Gln Lys 1 1835												5544
atg aca g Met Thr V 1850		aag gtc Lys Val			_		_			gac Asp		5589
tac aaa k Tyr Lys ( 1865												5634
gga ggc o Gly Gly I 1880	_				_					cct Pro	_	5679
aag gcc a Lys Ala 1 1895												5724
cgg gtt t Arg Val I 1910		cga caa Arg Gln	_	_	_				-	aat Asn		5769
ggc gcc a Gly Ala 1 1925												5814
tgg ggc a Trp Gly 1	Thr Gln		Gln	Asp	Val	Leu	Tyr	Glu	Āla			5859
cct cta c Pro Leu ( 1955												5904
gga gca c Gly Ala I 1970										ttg Leu	-	5949
gga ccg g Gly Pro <i>H</i> 1985												5994

	gcc Ala 2000		_		_	ctg Leu 2005		_	_		_	_	_		6039
	gcc Ala 2015	-		-		cgc Arg 2020					_	_		_	6084
_	cat His 2030					tgc Cys 2035									6129
	agc Ser 2045					cac His 2050									6174
	gac Asp 2060					tcc Ser 2065									6219
_	gac Asp 2075		_		_	cca Pro 2080	_		_	_				_	6264
	cat His 2090					aca Thr 2095									6309
	cga Arg 2105					gag Glu 2110							tct Ser		6354
	gag Glu 2120					tat Tyr 2125							agc Ser		6399
	ccc Pro 2135				_	ccc Pro 2140			_	_			ata Ile	_	6444
	aca Thr 2150					cca Pro 2155							agt Ser		6489
						ttg Leu 2170								acg Thr	6534
_	ggc Gly 2180					agg Arg 2185									6579
			_			tac Tyr 2200	_	_	_			_		_	6624

	ttt Phe 2210				agt Ser 2215							6669
_	ctc Leu 2225	_	_		tct Ser 2230	_		_	_	_		6714
	ccc Pro 2240	_	_	_	cta Leu 2245	_			_			6759
					cgt Arg 2260							6804
	ctg Leu 2270				ctc Leu 2275							6849
	tct Ser 2285				agg Arg 2290							6894
	caa Gln 2300				cgc Arg 2305							6939
					ggc Gly 2320							6984
	cca Pro 2330				tgg Trp 2335		tag					7011

<210> 30

<211> 2336

<212> PRT

<213> Rattus norvegicus

<400> 30

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Thr Gly 1 5 10 15

Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro
20 25 30

Gly Gln Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80

55

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140

Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser 145 150 155 160

Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Leu Thr 165 170 175

Glu Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 185 190

Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205

Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 215 220

Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile 225 230 235 240

Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 255

Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala 260 265 270

Pro Ala Arg Leu Cys Asp Ser Asp Thr Glu Cys Arg Glu Tyr Trp Pro 275 280 285

Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 290 295 300

Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 305 310 315 320

Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 325 330 335

Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 350

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365

Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 380

Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400

Ala Glu Glu Asp Lys Asn Ala Glu Glu Lys Ser Pro Leu Asp Ala Val 405 410 415

Leu Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala 420 425 430

Glu Glu Gly Glu Asp Arg Phe Val Asp Leu Cys Ala Ala Gly Ser Pro 435 440 445

Phe Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr 450 455 460

Phe Arg Arg Lys Glu Lys Met Phe Arg Phe Leu Ile Arg Arg Met Val 465 470 475 480

Lys Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn 485 490 495

Thr Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Gln Arg Leu Thr
500 505 510

- Thr Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr 515 520 525
- Glu Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg 530 535 540
- Ser Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe 545 550 555
- Glu Val Val Trp Ala Ala Ile Lys Pro Gly Thr Ser Phe Gly Ile Ser 565 570 575
- Val Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr 580 585 590
- Trp Asn Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys 595 600 605
- Ser Ile Ile Ser Leu Leu Phe Leu Phe Leu Phe Ile Val Val Phe 610 615 620
- Ala Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp 625 630 635 640
- Glu Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr 645 650 655
- Val Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His 660 665 670
- Gly Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe 675 680 685
- Tyr Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val 690 695 700
- Phe Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr 705 710 715 720
- Lys Asp Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu 725 730 735

- Gln Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn 740 745 750
- Ile Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val 755 760 765
- Trp Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser 770 780
- Cys Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Tyr 785 790 795 800
- Ala Ser Thr Arg His Val Arg Pro Asp Met Lys Thr His Met Asp Arg 805 810 815
- Pro Leu Val Val Glu Pro Gly Arg Asp Gly Leu Arg Gly Pro Ala Gly 820 825 830
- Asn Lys Ser Lys Pro Glu Gly Thr Glu Ala Thr Glu Gly Ala Asp Pro 835 840 845
- Pro Arg Arg His His Arg His Arg Asp Arg Asp Lys Thr Ser Ala Ser 850 860
- Thr Pro Ala Gly Glu Gln Asp Arg Thr Asp Cys Pro Lys Ala Glu 865 870 875 880
- Ser Thr Glu Thr Gly Ala Arg Glu Glu Arg Ala Arg Pro Arg Arg Ser 885 890 895
- His Ser Lys Glu Ala Pro Gly Ala Asp Thr Gln Val Arg Cys Glu Arg 900 905 910
- Ser Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Thr Glu Arg 915 920 925
- Glu Pro Arg Arg His Arg Ala His Arg His Ala Gln Asp Ser Ser Lys 930 935 940
- Glu Gly Lys Glu Gly Thr Ala Pro Val Leu Val Pro Lys Gly Glu Arg 945 950 955 960
- Arg Ala Arg His Arg Gly Pro Arg Thr Gly Pro Arg Glu Thr Glu Asn

965 970 975

- Ser Glu Glu Pro Thr Arg Arg His Arg Ala Lys His Lys Val Pro Pro 980 985 990
- Thr Leu Glu Pro Pro Glu Arg Glu Val Ala Glu Lys Glu Ser Asn Val 995 1000 1005
- Val Glu Gly Asp Lys Glu Thr Arg Asn His Gln Pro Lys Glu Pro 1010 1015 1020
- Arg Cys Asp Leu Glu Ala Ile Ala Val Thr Gly Val Gly Ser Leu 1025 1030 1035
- His Met Leu Pro Ser Thr Cys Leu Gln Lys Val Asp Glu Gln Pro 1040 1045 1050
- Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly Ser Gln 1055 1060 1065
- Pro Ser Asp Pro Ser Thr Thr Val His Val Pro Val Thr Leu Thr 1070 1075 1080
- Gly Pro Pro Gly Glu Ala Thr Val Val Pro Ser Ala Asn Thr Asp 1085 1090 1095
- Leu Glu Gly Gln Ala Glu Gly Lys Lys Glu Ala Glu Ala Asp Asp 1100 1105 1110
- Val Leu Arg Arg Gly Pro Arg Pro Ile Val Pro Tyr Ser Ser Met 1115 1120 1125
- Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys His Tyr 1130 1140
- Ile Val Thr Met Arg Tyr Phe Glu Met Val Ile Leu Val Val Ile 1145 1150 1155
- Ala Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val Arg Thr 1160 1165 1170
- Asp Ser Phe Arg Asn Asn Ala Leu Lys Tyr Met Asp Tyr Ile Phe 1175 1180 1185

- Thr Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile Asp Leu 1190 1195 1200
- Gly Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu Trp Asn 1205 1210 1215
- Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala Phe Ala 1220 1225 1230
- Phe Ser Ser Phe Met Gly Gly Ser Lys Gly Lys Asp Ile Asn Thr 1235 1240 1245
- Ile Lys Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr 1250 1260
- Ile Lys Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val 1265 1270 1275
- Asn Ser Leu Lys Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu 1280 1285 1290
- Phe Met Phe Ile Phe Ala Val Ile Ala Val Gln Leu Phe Lys Gly 1295 1300 1305
- Lys Phe Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp 1310 1315 1320
- Cys Arg Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala 1325 1330 1335
- Gln Pro Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr Asp Asn Val 1340 1345 1350
- Leu Trp Ala Leu Leu Thr Leu Phe Thr Val Ser Thr Gly Glu Gly 1355 1360 1365
- Trp Pro Met Val Leu Lys His Ser Val Asp Ala Thr Tyr Glu Glu 1370 1375 1380
- Gln Gly Pro Ser Pro Gly Phe Arg Met Glu Leu Ser Ile Phe Tyr 1385 1390 1395

- Val Val Tyr Phe Val Val Phe Pro Phe Phe Phe Val Asn Ile Phe 1400 1405 1410
- Val Ala Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp Lys Val 1415 1420 1425
- Met Ser Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp 1430 1435 1440
- Phe Ala Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn 1445 1450 1455
- Lys Gln Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro 1460 1465 1470
- Pro Phe Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val 1475 1480 1485
- Val Leu Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu 1490 1495 1500
- Met Leu Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Leu 1505 1510 1515
- Glu Cys Ile Leu Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe 1520 1530
- Arg Asp Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly Ser 1535 1540 1545
- Ile Thr Asp Ile Leu Val Thr Glu Ile Ala Asn Asn Phe Ile Asn 1550 1560
- Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu 1565 1570 1575
- Cys Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val 1580 1590
- Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu Ile Ala Met 1595 1600 1605

- Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val Phe Gly Asn 1610 1615 1620
- Ile Ala Leu Asp Asp Gly Thr Ser Ile Asn Arg His Asn Asn Phe 1625 1630 1635
- Arg Thr Phe Leu Gln Ala Leu Met Leu Leu Phe Arg Ser Ala Thr 1640 1645 1650
- Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys Leu Gly Asn Arg 1655 1660 1665
- Ala Cys Asp Pro His Ala Asn Ala Ser Glu Cys Gly Ser Asp Phe 1670 1680
- Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser Phe Leu 1685 1690 1695
- Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu Tyr 1700 1705 1710
- Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu 1715 1720 1725
- Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys Gly Arg 1730 1740
- Ile Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His Met Ser Pro 1745 1750 1755
- Pro Leu Gly Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr Lys 1760 1765 1770
- Arg Leu Val Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr 1775 1780 1785
- Val His Phe Thr Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu 1790 1795 1800
- Glu Ile Lys Leu Ala Pro Ala Gly Thr Lys Gln His Gln Cys Asp 1805 1810 1815
- Ala Glu Leu Arg Lys Glu Ile Ser Ser Val Trp Ala Asn Leu Pro

1820 1825 1830

Gln	Lys 1835	Thr	Leu	Asp	Leu	Leu 1840	Val	Pro	Pro	His	Lys 1845	Pro	Asp	Glu
Met	Thr 1850	Val	Gly	Lys	Val	Tyr 1855	Ala	Ala	Leu	Met	Ile 1860	Phe	Asp	Phe
Tyr	Lys 1865	Gln	Asn	Lys	Thr	Thr 1870	Arg	Asp	Gln	Thr	His 1875	Gln	Ala	Pro
Gly	Gly 1880	Leu	Ser	Gln	Met	Gly 1885	Pro	Val	Ser	Leu	Phe 1890	His	Pro	Leu
Lys	Ala 1895	Thr	Leu	Glu	Gln	Thr 1900	Gln	Pro	Ala	Val	Leu 1905	Arg	Gly	Ala
Arg	Val 1910	Phe	Leu	Arg	Gln	Lys 1915	Ser	Ala	Thr	Ser	Leu 1920	Ser	Asn	Gly
Gly	Ala 1925	Ile	Gln	Thr	Gln	Glu 1930	Ser	Gly	Ile	Lys	Glu 1935		Leu	Ser
Trp	Gly 1940		Gln	Arg	Thr	Gln 1945	Asp	Val	Leu	Tyr	Glu 1950	Ala	Arg	Ala
Pro	Leu 1955	Glu	Arg	Gly	His	Ser 1960	Ala	Glu	Ile	Pro	Val 1965	Gly	Gln	Pro
Gly	Ala 1970	Leu	Ala	Val	Asp	Val 1975	Gln	Met	Gln	Asn	Met 1980	Thr	Leu	Arg
Gly	Pro 1985	Asp	Gly	Glu	Pro	Gln 1990	Pro	Gly	Leu	Glu	Ser 1995	Gln	Gly	Arg
Ala	Ala 2000	Ser	Met	Pro	Arg	Leu 2005	Ala	Ala	Glu	Thr	Gln 2010	Pro	Ala	Pro
Asn	Ala 2015	Ser	Pro	Met	Lys	Arg 2020	Ser	Ile	Ser	Thr	Leu 2025	Ala	Pro	Arg
Pro	His 2030	Gly	Thr	Gln	Leu	Cys 2035	Asn	Thr	Val	Leu	Asp 2040	Arg	Pro	Pro

- Pro Ser Gln Val Ser His His His His His Arg Cys His Arg Arg 2045 2050 2055
- Arg Asp Lys Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu Ser 2060 2065 2070
- Val Asp Thr Glu Gly Ala Pro Ser Thr Ala Ala Gly Ser Gly Leu 2075 2080 2085
- Pro His Gly Glu Gly Ser Thr Gly Cys Arg Arg Glu Arg Lys Gln 2090 2095 2100
- Glu Arg Gly Arg Ser Gln Glu Arg Arg Gln Pro Ser Ser Ser Ser 2105 2110 2115
- Ser Glu Lys Gln Arg Phe Tyr Ser Cys Asp Arg Phe Gly Ser Arg 2120 2125 2130
- Glu Pro Pro Gln Pro Lys Pro Ser Leu Ser Ser His Pro Ile Ser 2135 2140 2145
- Pro Thr Ala Ala Leu Glu Pro Gly Pro His Pro Gln Gly Ser Gly 2150 2155 2160
- Ser Val Asn Gly Ser Pro Leu Met Ser Thr Ser Gly Ala Ser Thr 2165 2170 2175
- Pro Gly Arg Gly Gly Arg Arg Gln Leu Pro Gln Thr Pro Leu Thr 2180 2185 2190
- Pro Arg Pro Ser Ile Thr Tyr Lys Thr Ala Asn Ser Ser Pro Val 2195 2200 2205
- His Phe Ala Glu Gly Gln Ser Gly Leu Pro Ala Phe Ser Pro Gly 2210 2215 2220
- Arg Leu Ser Arg Gly Leu Ser Glu His Asn Ala Leu Leu Gln Lys 2225 2230 2235
- Glu Pro Leu Ser Gln Pro Leu Ala Ser Gly Ser Arg Ile Gly Ser 2240 2245 2250

Asp	2255	ıyı	пец	Gly	GIII	2260		ı Ası	, se	:I G		2265	ser	Ala	uis			
Asn	Leu 2270	Pro	Glu	Asp	Thr	Leu 2275		c Phe	e Gl	.u Gl		Ala 2280	Val	Ala	Thr			
Asn	Ser 2285	Gly	Arg	Ser	Ser	Arg 2290		s Sei	с Ту	r Vá		Ser 2295	Ser	Leu	Thr			
Ser	Gln 2300	Ser	His	Pro	Leu	Arg 2305	-	g Vai	l Pr	o As		Gly 2310	Tyr	His	Cys			
Thr	Leu 2315	Gly	Leu	Ser	Thr	Gly 2320		l Arg	g Al	.a Aı	_	His 2325	Ser	Tyr	His			
His	Pro 2330	Asp	Gln	. Asp	His	Trp 2335	-	5										
<213 <212	0> 33 L> 73 2> Di B> Ho	364 NA	sapi	ens														
<220 <220 <220	L> CI		(7	165)														
<400 gcgg			tgcg	gcgg	t ggg	ggccg	ggc	gag	gtcc	gct	gcg	ggtco	cgg	cggc	tccgtg	ī	60	
gct	gataco	gc t	ctga	gcgc	c tg	gegeg	ccc	cgc	gccc	tcc	cto	gccgg	ggc	cgct	gggccg	Ī	120	
ggga	atgcad	eg e	g <b>g</b> gg	cccg	g gag		et 1								ggc Gly		172	
	cgc t Arg T			Gly													220	
	ggc g Gly (							Pro (							cag Gln		268	
	gtc o Val I	eu					le A										316	
ctg	tac a	aac	ccc	atc	ccg q	gtc a	ag d	cag a	aac	tgc	tto	acc	gto	aac	cgc		364	

Leu	Tyr	Asn 60	Pro	Ile	Pro	Val	Lys 65	Gln	Asn	Cys	Phe	Thr 70	Val	Asn	Arg	
							gac Asp									412
							ttc Phe									460
							gcc Ala									508
							cgg Arg									556
							gca Ala 145									604
							tac Tyr									652
							ggg Gly									700
ttc Phe	gac Asp	ctg Leu	cga Arg	aca Thr 190	ctg Leu	agg Arg	gct Ala	gtg Val	cgt Arg 195	gtg Val	ctg Leu	agg Arg	ccc Pro	ctg Leu 200	aag Lys	748
_						_	ttg Leu	_				_			_	796
_	_	_	_			_	cag Gln 225			_					_	844
							ggc Gly									892
							agc Ser		_						_	940
							cca Pro									988
	_		_				gga Gly									1036

285 290 295

_			_		-		ttg Leu 305	_			_	_			_	108	34
_ ~				_			tat Tyr				_	~ ~	_			113	32
							atc Ile									118	30
	_			_		_	ggc Gly			_				_	_	122	28
							cgc Arg	-	_		_	_	_	_		127	76
_	_	_			_		ctc Leu 385				_					132	24
							gcc Ala									137	72
			_	_		_	aag Lys	_		_		_	_	_	_	142	20
							gag Glu									146	58
							gcc Ala									151	۱6
							cgg Arg 465									156	54
							gct Ala									161	L 2
							ctg Leu									166	50
_	_					_	acc Thr	_			_			_		170	8(

					aca Thr											1756
	_	_			cgg Arg					_		_			_	1804
atc Ile	gtg Val 555	G1A aaa	agc Ser	gtc Val	ttt Phe	gaa Glu 560	gtg Val	gtc Val	tgg Trp	gcg Ala	gcc Ala 565	atc Ile	aag Lys	ccg Pro	gga Gly	1852
	Ser				agt Ser 575											1900
					tac Tyr											1948
_	_			_	aag Lys				_	_			_			1996
					ttc Phe											2044
					gat Asp											2092
					act Thr 655											2140
					cac His											2188
					ttt Phe											2236
					gtc Val											2284
				_	acc Thr	_	-	-			_	_	-	_	_	2332
					ctg Leu 735											2380

ccc at Pro Me															2428
gcc aa Ala Ly															2476
cag aa Gln As															2524
gag ga Glu Gl 79	u Arg	_	_		-		_	_		_			_	_	2572
aag ac Lys Th 810															2620
gcg cg Ala Ar							_	_			_			_	2668
ccc ga Pro Gl		_	_		_	_						_	_	_	2716
gac aa Asp Ly	-					_	_	_	_	-		_	_	_	2764
gcg ga Ala Gl 87	u Ser				-	-			_		_		_		2812
cgc ag Arg Se 890					-				_				_		2860
cgc gg Arg Gl	_														2908
ggc to Gly Se															2956
cac cg His Ar	_	_	_	_	_	_		_	_		_	_			3004
cgg cg Arg Ar 95	g Ala														3052
gag ag	jc g <u>g</u> g	gag	gag	ccg	gcg	cgg	cgg	cac	cgg	gcc	cgg	cac	aag	gcg	3100

Glu Ser Gly Glu 970	Glu Pro Ala Arg 975	Arg His Arg Ala Arg 980	His Lys Ala 985
		aag gag acc acg gag Lys Glu Thr Thr Glu 995	
	Ala Glu Ile Val	gaa gcc gac aag ga Glu Ala Asp Lys Gl 1010	
	Gln Pro Arg Glu	cca cac tgt gac ct Pro His Cys Asp Le 1025	
	Thr Val Gly Pro	atg cac aca ctg co Met His Thr Leu Pr 1040	
	Val Glu Glu Gln	cca gag gat gca ga Pro Glu Asp Ala As 1055	
	Arg Met Gly Ser	cag ccc cca gac co Gln Pro Pro Asp Pr 1070	<del></del>
	Pro Val Met Leu	acg ggc cct ctt gg Thr Gly Pro Leu Gl 1085	
	Ser Gly Asn Val	gac ctg gaa agc ca Asp Leu Glu Ser Gl 1100	
	Val Glu Ala Asp	gac gtg atg agg ag Asp Val Met Arg Se 1115	
	Pro Tyr Ser Ser	atg ttc tgt tta ag Met Phe Cys Leu Se 1130	
	Arg Phe Cys His	tac atc gtg acc at Tyr Ile Val Thr Me 1145	
ttc gag gtg gtc Phe Glu Val Val 1155	Ile Leu Val Val	atc gcc ttg agc ag Ile Ala Leu Ser Se 1160	
	Asp Pro Val Arg	aca gac tcg ccc ac Thr Asp Ser Pro Ar 1175	
_	ctg gat tac att Leu Asp Tyr Ile	, _	

		1185				1190					1195		
						ttg Leu 1205							3778
	_			_	_	 aac Asn 1220		_	_			gtg Val	3823
_	~	 -	_	~ ~		gct Ala 1235						gjå aaa	3868
	_				_	ctg Leu 1250	_	_		_	_	_	3913
						ctg Leu 1265							3958
						aag Lys 1280							4003
						ata Ile 1295							4048
						tac Tyr 1310						aag Lys	4093
	_	 	_			 cag Gln 1325		_	_			aag Lys	4138
	_	 _	_	_		 cag Gln 1340		_			_	ttt Phe	4183
						ctg Leu 1355						gtg Val	4228
						gtg Val 1370						gat Asp	4273
						agc Ser 1385							4318
_					_	ttt Phe 1400		_				ttc Phe	4363

ttc Phe	gtc Val	aac Asn	atc Ile 1410	ttt Phe	gtg Val	gct Ala	ttg Leu	atc Ile 1415	atc Ile	atc Ile	acc Thr	ttc Phe	cag Gln 1420	gag Glu	4408
_		_	aag Lys 1425		_		_	_	_	_		_		gag Glu	4453
			att Ile 1440											cgg Arg	4498
			caa Gln 1455											aca Thr	4543
			tcc Ser 1470											ata Ile	4588
Ala	Leu	Asn	act Thr 1485	Val	Val	Leu	Met	Met 1490	Lys	Phe	Tyr	Asp	Ala 1495	ccc Pro	4633
Tyr	Glu	Tyr	gag Glu 1500	Leu	Met	Leu	Lys	Cys 1505	Leu	Asn	Ile	Val	Phe 1510	aca Thr	4678
Ser	Met	Phe	tcc Ser 1515	Met	Glu	Cys	Val	Leu 1520	Lys	Ile	Ile	Āla	Phe 1525	_	4723
Val	Leu	Asn	tat Tyr 1530	Phe	Arg	Asp	Ala	Trp 1535	Asn	Val	Phe	Asp	Phe 1540	Val	4768
Thr	Val	Leu	gga Gly 1545	Ser	Ile	Thr	Asp	Ile 1550	Leu	Val	Thr	Glu	Ile 1555	Ala	4813
Glu	Thr	Asn	aat Asn 1560 ctg	Phe	Ile	Asn	Leu	Ser 1565	Phe	Leu	Arg	Leu	Phe 1570	Arg	4858
Ala	Ala	Arg	Leu 1575	I·le	Lys	Leu	Leu	Arg 1580	Gln	Gly	Tyr	Thr	Ile 1585	Arg	4903
Ile	Leu	Leu	tgg Trp 1590	Thr	Phe	Val	Gln	Ser 1595	Phe	Lys	Ala	Leu	Pro 1600	tac Tyr	4948
_	_	_	ctc Leu 1605		_	_	_					_		atc Ile	4993

•

ggc atg c Gly Met G											agc Ser	5038
atc aac c Ile Asn A	_				_		_		_	_	atg Met	5083
ctg ctg t Leu Leu P		_				_					atg Met	5128
ctg tcc t Leu Ser C											gcc Ala	5173
acc gag t Thr Glu C											ttc Phe	5218
atc ttc c Ile Phe L			_	_	_					_	gtg Val	5263
atc atg g Ile Met A											cta Leu	5308
ggt cct c Gly Pro H											tac Tyr	5353
gac ccg g Asp Pro A											gag Glu	5398
atg ctg a Met Leu L											tgc Cys	5443
cct gct c Pro Ala A	~ ~	_	r Lys	Arg	Leu	Val.	_	Met	Asn	_	ccc Pro	5488
atc tcc a Ile Ser A		_	_	_			_		_	_	_	5533
gcc ctc a Ala Leu I		acg gc Thr Al									gjå aaa	5578
aca aag c Thr Lys G	-		_			_					t.cc Ser	5623
gtt gtg t	.gg gcc	aat ct	g ccc	cag	aag.	act	ttg	gac	ttg	ctg	gta	5668

Val	Val	Trp	Ala 1830	Asn	Leu	Pro	Gln	Lys 1835	Thr	Leu	Asp	Leu	Leu 1840	Val	
			aag Lys 1845											gca Ala	5713
			ata Ile 1860												5758
			cag Gln 1875											cct Pro	5803
		_	ttc Phe 1890			_	_	-		_				cag Gln	5848
		-	ctc Leu 1905			-		_			_	_	_	agt Ser	5893
			ctc Leu 1920	_				-					gag Glu 1930	agt Ser	5938
			gag Glu 1935												5983
_			gag Glu 1950	_				_		_				aca Thr	6028
			gtg Val 1965											cag Gln	6073
			ata Ile 1980					cct Pro 1985			-		_	cct Pro	6118
			agc Ser 1995											gcg Ala	6163
			cag Gln 2010											tcc Ser	6208
			ctg Leu 2025											agc Ser	6253
		_	gac Asp	_				_	-		_	_		cac His	6298

2040	2045		2050	
cgc tgc cac cgc Arg Cys His Arg 2055		p Arg Lys Gln		6343
ggg ccc agc ctg Gly Pro Ser Leu 2070		p Met Asp Gly .		6388
gtg ggg ccg ggg Val Gly Pro Gly 2085		o Gly Glu Gly		6433
 cgg gaa cga gag Arg Glu Arg Glu 2100		n Glu Arg Gly		6478
agg cag ccc tca Arg Gln Pro Ser 2115		r Ser Glu Lys		6523
tgc gac cgc ttt Cys Asp Arg Phe 2130		g Glu Pro Pro		6568
ctc agc agc cac Leu Ser Ser His 2145		r Pro Thr Ala		6613
ccc cac cca cag Pro His Pro Gln 2160		y Ser Val Asn		6658
tca aca tct ggt Ser Thr Ser Gly 2175		r Pro Gly Arg		6703
 ctc ccc cag acg Leu Pro Gln Thr 2190		r Pro Arg Pro		6748
acg gcc aac tcc Thr Ala Asn Ser 2205		e His Phe Ala		6793
ctc cct gcc ttc Leu Pro Ala Phe 2220		y Arg Leu Ser .		6838
cac aac gcc ctg His Asn Ala Leu 2235		g Asp Pro Leu		6883
cct ggc tct cga Pro Gly Ser Arg 2250		r Asp Pro Tyr		6928

_	_	_	gac Asp 2265					gtc Val 2270						gac Asp	6973
_			ttc Phe 2280			_		gcc Ala 2285			_		_	tcc Ser	7018
								ctg Leu 2300			_			cct Pro	7063
								cac His 2315						agc Ser	7108
			_	_			_	tac Tyr 2330				_		gac Asp	7153
cac tgg tgc tag ctgcaccgtg accgctcaga cgcctgcatg cagcaggcgt 7 His Trp Cys															7205
gtgttccagt ggatgagttt tatcatccac acggggcagt cggccctcgg gggaggcctt														aggcctt	7265
geceacettg gtgaggetee tgtggeeeet eeeteeeet ecteeeetet tttactetag														actctag	7325
acga	acgacgaata aagccctgtt gcttgagtgt acgtaccgc														7364
<210> 32 <211> 2339															

<212> PRT

<213> Homo sapiens

<400> 32

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly 10

Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro 20 25 30

Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 70 75

- Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95
- Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110
- Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125
- Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140
- Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser 145 150 155 160
- Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Leu Thr 165 170 175
- Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 185 190
- Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205
- Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 215 220
- Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile 225 230 235 240
- Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 255
- Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala 260 265 270
- Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Pro 275 280 285
- Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 290 295 300

Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 305 310 315 320

Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 325 330 335

Ile Pro Leu Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 350

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365

Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 375 380

Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400

Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu 405 410 415

Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu 420 425 430

Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe 435 440 445

Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe 450 455 460

Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys 465 470 475 480

Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr 485 490 495

Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Arg Arg Leu Thr Thr 500 505 510

Thr Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 525

Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser 530 540

Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Val Phe Glu 545 550 555 560

Val Val Trp Ala Ala Ile Lys Pro Gly Ser Ser Phe Gly Ile Ser Val 565 570 575

Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp 580 585 590

Ser Ser Leú Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 595 600 605

Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala 610 615 620

Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 635 640

Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val 645 . 650 655

Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly 660 665 670

Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr 675 680 685

Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe 690 695 700

Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys 705 710 715 720

Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln 725 730 735

Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750

Ser Ile Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp

755 760 765

Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys 770 780

Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala 785 790 795 800

Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro 805 810 815

Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly 820 825 830

Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro 835 840 845

Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly 850 855

Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly 865 870 875 880

Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala 885 890 895

Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro 900 905 910

Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala 915 920 925

Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser 930 935 940

Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly 945 955 960

Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala 965 970 975

Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val 980 985 990

- Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile 995 1000 1005
- Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg 1010 1015 1020
- Glu Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly 1025 1030 1035
- Pro Met His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu 1040 1045 1050
- Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly 1055 1060 1065
- Ser Gln Pro Pro Asp Pro Asn Thr Ile Val His Ile Pro Val Met 1070 1075 1080
- Leu Thr Gly Pro Leu Gly Glu Ala Thr Val Val Pro Ser Gly Asn 1085 1090 1095
- Val Asp Leu Glu Ser Gln Ala Glu Gly Lys Lys Glu Val Glu Ala 1100 1105 1110
- Asp Asp Val Met Arg Ser Gly Pro Arg Pro Ile Val Pro Tyr Ser 1115 1120 1125
- Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys 1130 1135 1140
- His Tyr Ile Val Thr Met Arg Tyr Phe Glu Val Val Ile Leu Val 1145 1150 1155
- Val Ile Ala Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val 1160 1165 1170
- Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys Tyr Leu Asp Tyr 1175 1180 1185
- Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile 1190 1195 1200

- Asp Leu Gly Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu 1205 1210 1215
- Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala 1220 1225 1230
- Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys 1235 1240 1245
- Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys 1250 1255 1260
- Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser 1265 1270 1275
- Leu Lys Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met 1280 1285 1290
- Phe Ile Phe Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe 1295 1300 1305
- Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg 1310 1315 1320
- Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro 1325 1330 1335
- Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr Asp Asn Val Leu Trp 1340 1345 1350
- Ala Leu Leu Thr Leu Phe Thr Val Ser Thr Gly Glu Gly Trp Pro 1355 1360 1365
- Met Val Leu Lys His Ser Val Asp Ala Thr Tyr Glu Glu Gln Gly 1370 1375 1380
- Pro Ser Pro Gly Tyr Arg Met Glu Leu Ser Ile Phe Tyr Val Val 1385 1390 1395
- Tyr Phe Val Val Phe Pro Phe Phe Val Asn Ile Phe Val Ala 1400 1405 1410

- Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp Lys Val Met Ser 1415 1420 1425
- Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp Phe Ala 1430 1435 1440
- Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn Arg Gln 1445 1450 1455
- Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro Phe 1460 1465 1470
- Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu 1490 1495 1500
- Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys 1505 1510 1515
- Val Leu Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp 1520 1530
- Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr 1535 1540 1545
- Asp Ile Leu Val Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn 1550 1560
- Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu 1565 1570 1575
- Leu Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val 1580 1585 1590
- Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu Ile Ala Met 1595 1600 1605
- Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val Phe Gly Asn 1610 1615
- Ile Ala Leu Asp Asp Asp Thr Ser Ile Asn Arg His Asn Asn Phe

1625 1630 1635

Arg	Thr 1640	Phe	Leu	Gln	Ala	Leu 1645	Met	Leu	Leu	Phe	Arg 1650	Ser	Ala	Thr
Gly	Glu 1655	Ala	Trp	His	Glu	Ile 1660	Met	Leu	Ser	Cys	Leu 1665	Ser	Asn	Gln
Ala	Cys 1670	Asp	Glu	Gln	Ala	Asn 1675	Ala	Thr	Glu	Cys	Gly 1680	Ser	Asp	Phe
Ala	Tyr 1685	Phe	Tyr	Phe	Val	Ser 1690	Phe	Ile	Phe	Leu	Cys 1695	Ser	Phe	Leu
Met	Leu 1700	Asn	Leu	Phe	Val	Ala 1705	Val	Ile	Met	Asp	Asn 1710	Phe	Glu	Tyr
Leu	Thr 1715	Arg	Asp	Ser	Ser	Ile 1720	Leu	Gly	Pro	His	His 1725	Leu	Asp	Glu
Phe	Ile 1730	Arg	Val	Trp	Ala	Glu 1735	Tyr	Asp	Pro	Ala	Ala 1740	Cys	Gly	Arg
Ile	Ser 1745	Tyr	Asn	Asp	Met	Phe 1750	Glu	Met	Leu	Lys	His 1755	Met	Ser	Pro
Pro	Leu 1760	Gly	Leu	Gly		Lys 1765	Cys	Pro	Ala	Arg	Val 1770	Ala	Tyr	Lys
Arg	Leu 1775	Val	Arg	Met	Asn	Met 1780		Ile	Ser	Asn	Glu 1785	Asp	Met	Thr
Val	His 1790	Phe	Thr	Ser	Thr	Leu 1795	Met	Ala	Leu	Ile	Arg 1800	Thr	Ala	Leu
Glu	Ile 1805	Lys	Leu	Ala	Pro	Ala 1810	Gly	Thr	Lys	Gln	His 1815	Gln	Cys	Asp
Ala	Glu 1820	Leu	Arg	Lys	Glu	Ile 1825	Ser	Val	Val	Ťrp	Ala 1830	Asn	Leu	Pro
Gln	Lys 1835	Thr	Leu	Asp	Leu	Leu 1840	Val	Pro	Pro	His	Lys 1845	Pro	Asp	Glu

- Met Thr Val Gly Lys Val Tyr Ala Ala Leu Met Ile Phe Asp Phe 1850 1860
- Tyr Lys Gln Asn Lys Thr Thr Arg Asp Gln Met Gln Gln Ala Pro 1865 1870 1875
- Gly Gly Leu Ser Gln Met Gly Pro Val Ser Leu Phe His Pro Leu 1880 1885 1890
- Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Val Leu Arg Gly Ala 1895 1900 1905
- Arg Val Phe Leu Arg Gln Lys Ser Ser Thr Ser Leu Ser Asn Gly 1910 1915 1920
- Gly Ala Ile Gln Asn Gln Glu Ser Gly Ile Lys Glu Ser Val Ser 1925 1930 1935
- Trp Gly Thr Gln Arg Thr Gln Asp Ala Pro His Glu Ala Arg Pro 1940 1945 1950
- Pro Leu Glu Arg Gly His Ser Thr Glu Ile Pro Val Gly Arg Ser 1955 1960 1965
- Gly Ala Leu Ala Val Asp Val Gln Met Gln Ser Ile Thr Arg Arg 1970 1980
- Gly Pro Asp Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg 1985 1990 1995
- Ala Ala Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Val Thr 2000 2005 2010
- Asp Ala Ser Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Gln Arg 2015 2020 2025
- Pro Arg Gly Thr His Leu Cys Ser Thr Thr Pro Asp Arg Pro Pro 2030 2035 2040
- Pro Ser Gln Ala Ser Ser His His His His Arg Cys His Arg 2045 2050 2055

- Arg Arg Asp Arg Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu 2060 2065 2070
- Ser Ala Asp Met Asp Gly Ala Pro Ser Ser Ala Val Gly Pro Gly 2075 2080 2085
- Leu Pro Pro Gly Glu Gly Pro Thr Gly Cys Arg Arg Glu Arg Glu 2090 2095 2100
- Arg Arg Gln Glu Arg Gly Arg Ser Gln Glu Arg Arg Gln Pro Ser 2105 2115
- Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr Ser Cys Asp Arg Phe 2120 2125 2130
- Gly Gly Arg Glu Pro Pro Lys Pro Lys Pro Ser Leu Ser Ser His 2135 2140 2145
- Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro His Pro Gln 2150 2155 2160
- Gly Ser Gly Ser Val Asn Gly Ser Pro Leu Leu Ser Thr Ser Gly 2165 2170 2175
- Ala Ser Thr Pro Gly Arg Gly Gly Arg Arg Gln Leu Pro Gln Thr 2180 2185 2190
- Pro Leu Thr Pro Arg Pro Ser Ile Thr Tyr Lys Thr Ala Asn Ser 2195 2200 2205
- Ser Pro Ile His Phe Ala Gly Ala Gln Thr Ser Leu Pro Ala Phe 2210 2215 2220
- Ser Pro Gly Arg Leu Ser Arg Gly Leu Ser Glu His Asn Ala Leu 2225 2230 2235
- Leu Gln Arg Asp Pro Leu Ser Gln Pro Leu Ala Pro Gly Ser Arg 2240 2250
- Ile Gly Ser Asp Pro Tyr Leu Gly Gln Arg Leu Asp Ser Glu Ala 2255 2260 2265

Ser	Val 2270		. Ala	. Leu	Pro	Glu 227		зр Т	hr 1	Leu	Th		he 280	Glu	Glu	Al	a	
Val	Ala 2285		Asn	ser	Gly	Arc 229		er S	er 1	Arg	Th		er 295	Tyr	Val	Se	er	
Ser	Leu 2300		Ser	Gln	ı Ser	His 230		ro L	eu i	Arg	Ar	_	al 310	Pro	Asn	Gl	У	
Туг	His 2315	-	Thr	Leu	ı Gly	Leu 232		er S	er (	Gly	Gl	-	rg 325	Ala	Arg	Hi	s	
Ser	Tyr 2330		; His	Pro	Asp	Glr 233		sp H	is ?	Гrр	Су	rs						
<210 <211 <212 <213	L> 7	3 177 NA Como	sapi	.ens												,		
<220 <220 <220	L> C	DS 146)	(6	5859)														
<400 gcg		3 3	etgeg	gege	jt gg	ggcc	:999	c ga	ggt	ccg	ct	gcg	gtco	cgg	cggo	ctc	cgtg	60
gct	getee	gc t	ctga	gcgc	c tg	gcgc	gcc	c cg	cgc	cct	cc	ctg	ccgo	gggc	cgct	tgg	gccg	120
ggg	atgca	.cg c	gggg	leceē	gg ga										g cto ı Lei			172
	cgc Arg										rg					уĞ		220
	ggc Gly																	268
	gtc Val			_		_					_			•	-			316
	tac Tyr																	364
	ctc Leu																	412

_		 	cca Pro				_		_	_		460
			ctg Leu									508
_	_	_	gag Glu		_	_	_	_				556
			gag Glu									604
			tct Ser 160									652
			aca Thr									700
			agg Arg									748
_			agt Ser	_	_				-		_	796
			ctg Leu	_			_				_	844
			att Ile 240									892
			aac Asn	_		_	_				 _	.940
			gcc Ala									988
			cca Pro									1036
			atc Ile									1084

									aat Asn								1132
7									cct Pro								1180
		_			_		_		gtg Val		_				_	_	1228
									cgc Arg 370								1276
	_	_	_			_			aac Asn			_					1324
									gag Glu								1372
I									aga Arg								1420
ē I	aat Asn	gac Asp	ctg Leu	atc Ile	cac His 430	gca Ala	gag Glu	gag Glu	gga Gly	gag Glu 435	gac Asp	cgg Arg	ttt Phe	gca Ala	gat Asp 440	ctc Leu	1468
									cgc Arg 450								1516
									agg Arg								1564
	Phe			-	_		_	_	cag Gln							-	1612
C									tgt Cys								1660
									ctg Leu								1708
	_							_	tcc Ser 530	_	_	_			_		1756

.

.

														Gly 939		1804
														ccg Pro		1852
_					_		_		_		_	_	_	agg Arg		1900
														gtg Val 600		1948
														ctc Leu		1996
														GJÀ aaa		2044
_				_	_								_	acc Thr		2092
														gac Asp		2140
														agc Ser 680		2188
														gga Gly		2236
														ctg Leu		2284
Asn	Ala 715	Gln	Glu	Leu	Thr	Lys 720	Āsp	Glu	Glu	Glu	Met 725	Glu ,	Glu	gca Ala	Ala	2332
Asn 730	Gln	Lys	Leu	Āla	Leu 735	Gln	Lys	Āla	Lys	Glu 740	Val	Āla	Glu	gtc Val	Ser 745	2380
	_		_						_	_		_	_	aac Asn 760	_	2428
gcc	aag.	gcg	cgc	tcg	gtg	tgg	gag	cag	cgg	gcc	agc	cag	cta	cgg	ctg	2476

Ala	Lys	Ala	Arg 765	Ser	Val	Trp	Glu	Gln 770	Arg	Ala	Ser	Gln	Leu 775	Arg	Leu		
												gag Glu 790				2524	
gag Glu												cgg Arg				2572	
aag Lys 810												ggc Gly				2620	
gcg Ala								_	_			gct Ala			_	2668	
			_	_		_	_					cac His	_	_	~	2716	
_	_						_	_	_	_	-	gag Glu 870	_	_	_	2764	
												ccg Pro				2812	
_	_		_	_						_		gcg Ala		_		2860	
_		_										cac His			_	2908	
												cgc Arg				2956	
						Ser						gcc Ala 950				3004	
												ccc Pro				3052	
												cgg Arg				3100	
															gcc Ala	3148	

990 995 1000

			gag Glu 1005					gaa Glu 1010					aag Lys 1015	gag Glu	3193
			cac His 1020					cca Pro 1025						acc Thr	3238
_			gtg Val 1035					atg Met 1040			_		_	acc Thr	3283
-		_	aag Lys 1050			-	_			_	_	_		cag Gln	3328
		-	act Thr 1065	_	_		-	cag Gln 1070			_	_		act Thr	3373
			atc Ile 1080										gaa Glu 1090		3418
_	_	_	ccc Pro 1095	_				gac Asp 1100	_		_		_	gag Glu	3463
			gag Glu 1110										ggc Gly 1120	ccc Pro	3508
			gtc Val 1125					atg Met 1130						acc Thr	3553
	_		cgc Arg 1140	_		-						_		tac Tyr	3598
			gtc Val 1155	Ile			Val	atc Ile 1160	Āla	_	_	_		gcc Ala	3643
			gag Glu 1170										aac Asn 1180	aac Asn	3688
			tac Tyr 1185										acc Thr 1195	ttt Phe	3733
			ata Ile 1200										cac His 1210	cct Pro	3778

	_		ttc Phe 1215	 _	_				_	_			3823
			gcc Ala 1230										3868
			aat Asn 1245									ctg Leu	3913
			aag Lys 1260									gtg Val	3958
			gtg Val 1275										4003
			atg Met 1290									gtg Val	4048
_			aaa Lys 1305	 _				_		_	_	aag Lys	4093
			agg Arg 1320				cag Gln 1325		_	_		 aag Lys	4138
			gaa Glu 1335									ttt Phe	4183
		_	aat Asn 1350			_	ctg Leu 1355	_	_	_		gtg Val	4228
	_		gaa Glu 1365				gtg Val 1370					gat Asp	4273
			gag Glu 1380									gag Glu	4318
			ttc Phe 1395									ttc Phe	4363
			atc Ile 1410									gag Glu	4408

Gln Gly Asp Ly			ctg gag aag aac Leu Glu Lys Asn 1435	*	3
Arg Ala Cys Il			e aaa ccc ctg aca Lys Pro Leu Thr 1450	cgg 4498 Arg	3
Tyr Met Pro G			tat aag acg tgg Tyr Lys Thr Trp 1465	aca 4543 Thr	3
Phe Val Val Se			e atc atg gcc atg e Ile Met Ala Met 1480	ata 4588 Ile	3
Ala Leu Asn Th			ttc tat gat gca Phe Tyr Asp Ala 1495	ccc 4633	3
Tyr Glu Tyr Gl			g aac atc gtg ttc Asn Ile Val Phe 1510	aca 4678 Thr	
Ser Met Phe Se			g atc atc gcc ttt s Ile Ile Ala Phe 1525	ggg 4723 Gly	3
Val Leu Asn Ty			gtc ttt gac ttt Val Phe Asp Phe 1540	Val	
Thr Val Leu Gl	-	_	gta aca gag att Val Thr Glu Ile 1555	Ala	3
Glu Thr Asn As	sn Phe Ile Asn 560	Leu Ser Phe 1565	ctc cgc ctc ttt Leu Arg Leu Phe 1570	Arg	3
Ala Ala Arg Le 15	eu Ile Lys Leu 575	Leu Arg Gln 1580	ggc tac acc atc Gly Tyr Thr Ile 1585	Arg	3
Ile Leu Leu Tr			aag gcc ctg ccc Lys Ala Leu Pro 1600	tac 4948 Tyr	3
Val Cys Leu Le			e atc tac gcc atc e Ile Tyr Ala Ile 1615	atc 4993 Ile	
Gly Met Gln Va			gat gat gac acc Asp Asp Asp Thr 1630	agc 5038 Ser	3
atc aac cgc ca	ac aac ttc	cgg acg ttt	ttg caa gcc ctg	atg 5083	3

:	Ile	Asn	Arg	His 1635	Asn	Asn	Phe	Arg	Thr 1640	Phe	Leu	Gln	Ala	Leu 1645	Met		
									gag Glu 1655							5128	
									tgt Cys 1670							5173	
									tac Tyr 1685							5218	
									ttg Leu 1700							5263	
		_	_						acg Thr 1715		_					5308	
									atc Ile 1730							5353	
									agt Ser 1745							5398	
									ctg Leu 1760							5443	
									ctg Leu 1775							5488	
									cac His 1790							5533	
									atc Ile 1805						gjå aaa	5578	
		_	_						gag Glu 1820						tcc Ser	5623	
									aag Lys 1835							5668	
				_		_		_	aca Thr			_	_		gca Ala	5713	

	1845	1850		1855	
gct ctg atg Ala Leu Met	ata ttt gac Ile Phe Asp 1860	ttc tac aag Phe Tyr Lys 1865	cag aac aaa ac Gln Asn Lys Th	c acc aga c Thr Arg 1870	5758
gac cag atg Asp Gln Met			ctc tcc cag ato Leu Ser Gln Me		5803
gtg tcc ctg Val Ser Leu		ctg aag gcc Leu Lys Ala 1895			5848
ccg gct gtg Pro Ala Val			ttc ctt cga cag Phe Leu Arg Gli		5893
tcc acc tcc Ser Thr Ser			ata caa aac caa Ile Gln Asn Gla		5938
ggc atc aaa Gly Ile Lys			act caa agg acc Thr Gln Arg Th		5983
gca ccc cat Ala Pro His			gag cgt ggc cae Glu Arg Gly His		6028
gag atc cct Glu Ile Pro			ctg gct gtg ga Leu Ala Val As	_	6073
atg cag agc Met Gln Ser		agg ggc cct Arg Gly Pro 1985	gat ggg gag cc Asp Gly Glu Pro	_	6118
ggg ctg gag Gly Leu Glu		cga gcg gcc Arg Ala Ala 2000	tcc atg ccc cg Ser Met Pro Arg	<b>J J</b>	6163
			agc ccc atg aag Ser Pro Met Ly		6208
			ggg act cat ct Gly Thr His Le		6253
			cag gcg tcg tcg Gln Ala Ser Se		6298
			gac agg aag cag Asp Arg Lys Gli		6343

ctg gag aag ggg ccc agc ctg tct gcc gat atg gat ggc gca cca Leu Glu Lys Gly 'Pro Ser Leu Ser Ala Asp Met Asp Gly Ala Pro 2070 2075 2080	6388
agc agt gct gtg ggg ccg ggg ctg ccc ccg gga gag ggg cct aca Ser Ser Ala Val Gly Pro Gly Leu Pro Pro Gly Glu Gly Pro Thr 2085 2090 2095	6433
ggc tgc cgg cgg gaa cga gag cgc cgg cag gag cgg ggc cgg tcc Gly Cys Arg Arg Glu Arg Glu Arg Gln Glu Arg Gly Arg Ser 2100 2105 2110	6478
cag gag cgg agg cag ccc tca tcc tcc tcc tcg gag aag cag cgc Gln Glu Arg Arg Gln Pro Ser Ser Ser Ser Glu Lys Gln Arg 2115 2120 2125	6523
ttc tac tcc tgc gac cgc ttt ggg ggc cgt gag ccc ccg aag ccc Phe Tyr Ser Cys Asp Arg Phe Gly Gly Arg Glu Pro Pro Lys Pro 2130 2135 2140	6568
aag ccc tcc ctc agc agc cac cca acg tcg cca aca gct ggc cag Lys Pro Ser Leu Ser Ser His Pro Thr Ser Pro Thr Ala Gly Gln 2145 2150 2155	6613
gag ccg gga ccc cac cca cag gcc ggc tca gcc gtg ggc ttt ccg Glu Pro Gly Pro His Pro Gln Ala Gly Ser Ala Val Gly Phe Pro 2160 2165 2170	6658
aac aca acg ccc tgc tgc aga gag acc ccc tca gcc agc ccc tgg Asn Thr Thr Pro Cys Cys Arg Glu Thr Pro Ser Ala Ser Pro Trp 2175 2180 2185	6703
ccc ctg gct ctc gaa ttg gct ctg acc ctt acc tgg ggc agc gtc Pro Leu Ala Leu Glu Leu Ala Leu Thr Leu Thr Trp Gly Ser Val 2190 2195 2200	6748
tgg aca gtg agg cct ctg tcc acg ccc tgc ctg agg aca cgc tca Trp Thr Val Arg Pro Leu Ser Thr Pro Cys Leu Arg Thr Arg Ser 2205 2210 2215	6793
ctt tcg agg agg ctg tgg cca cca act cgg gcc gct cct cca gga Leu Ser Arg Arg Leu Trp Pro Pro Thr Arg Ala Ala Pro Pro Gly 2220 2225 2230	6838
ctt cct acg tgt cct ccc tga cctcccagtc tcaccctctc cgccgcgtgc Leu Pro Thr Cys Pro Pro 2235	6889
ccaacggtta ccactgcacc ctgggactca gctcgggtgg ccgagcacgg cacagctacc	6949
accaccetga ccaagaceae tggtgetage tgeacegtga cegeteagae geetgeatge	7009
agcaggcgtg tgttccagtg gatgagtttt atcatccaca cggggcagtc ggccctcggg	7069
ggaggeettg cecaeettgg tgaggeteet gtggeeeete eeteeeeete eteeeetett	7129

<210> 34

<211> 2237 <212> PRT <213> Homo sapiens

<400> 34

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly 10

Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro

Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 120 115

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135

Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser 155 .

Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr 165

Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 185

Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met 220 Val Pro Leu Leu 215 Ile Gly Leu Glu Phe 230 Phe Phe Ala Ile 235 Met Phe Ala Ile 240 Phe 240 Phe 240 Phe 245 Phe 255 Phe 25

Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser

195

340

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365

Ile Pro Leu Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu

345

- Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 380
- Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400
- Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu \$405\$ \$410\$ \$415\$

Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu 420 425 430

Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe 435 440 445

Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe 450 455 460

Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys 465 470 475 480

Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr 485 490 495

Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Arg Arg Leu Thr Thr 500 505 510

Thr Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 525

Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser 530 540

Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Val Phe Glu 545 550 555 560

Val Val Trp Ala Ala Ile Lys Pro Gly Ser Ser Phe Gly Ile Ser Val 565 570 575

Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp
580 585 590

Ser Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 595 600 605

Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala 610 615 620

Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 635 640

Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val

645 650 655

Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly 660 665 670

- Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr 675 680 685
- Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe 690 695 700
- Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys
  705 710 715 720
- Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln 725 730 735
- Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750
- Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp  $755 \hspace{1.5cm} 760 \hspace{1.5cm} 765$
- Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys
  770 780
- Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala 785 790 795 800
- Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro 805 810 815
- Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly 820 825 830
- Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro 835 840 845
- Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly 850 855
- Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly 865 870 875 880

- Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala 885 890 895
- Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro 900 905 910
- Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala 915 920 925
- Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser 930 935 940
- Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly 945 950 955 960
- Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala 965 970 975
- Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val 980 985 990
- Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile 995 1000 1005
- Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg 1010 1015 1020
- Glu Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly 1025 1030 1035
- Pro Met His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu 1040 1045 1050
- Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly 1055 1060 1065
- Ser Gln Pro Pro Asp Pro Asn Thr Ile Val His Ile Pro Val Met 1070 1075 1080
- Leu Thr Gly Pro Leu Gly Glu Ala Thr Val Val Pro Ser Gly Asn 1085 1090 1095

- Val Asp Leu Glu Ser Gln Ala Glu Gly Lys Lys Glu Val Glu Ala 1100 1105 1110
- Asp Asp Val Met Arg Ser Gly Pro Arg Pro Ile Val Pro Tyr Ser 1115 1120 1125
- Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys 1130 1140
- His Tyr Ile Val Thr Met Arg Tyr Phe Glu Val Val Ile Leu Val 1145 1150 1155
- Val Ile Ala Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val 1160 1165 1170
- Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys Tyr Leu Asp Tyr 1175 1180 1185
- Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile 1190 1195 1200
- Asp Leu Gly Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu 1205 1210 1215
- Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala 1220 1230
- Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys 1235 1240 1245
- Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys 1250 1260
- Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser 1265 1270 1275
- Leu Lys Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met 1280 1285 1290
- Phe Ile Phe Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe 1295 1300 1305

- Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg 1310 1315 1320
- Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro 1325 1330 1335
- Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr Asp Asn Val Leu Trp 1340 1345 1350
- Ala Leu Leu Thr Leu Phe Thr Val Ser Thr Gly Glu Gly Trp Pro 1355 1360 1365
- Met Val Leu Lys His Ser Val Asp Ala Thr Tyr Glu Glu Gln Gly 1370 1375 1380
- Pro Ser Pro Gly Tyr Arg Met Glu Leu Ser Ile Phe Tyr Val Val 1385 1390 1395
- Tyr Phe Val Val Phe Pro Phe Phe Val Asn Ile Phe Val Ala 1400 1405 1410
- Leu Ile Ile Thr Phe Gln Glu Gln Gly Asp Lys Val Met Ser 1415 1420 1425
- Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp Phe Ala 1430 1440
- Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn Arg Gln 1445 . 1450 1455
- Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro Phe 1460 1465 1470
- Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu 1475 1480 1485
- Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu 1490 1495 1500
- Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys 1505 1510 1515
- Val Leu Lys ïle Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp

1520 1525 1530

Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr Asp Ile Leu Val Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu Leu Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val Phe Gly Asn Ile Ala Leu Asp Asp Asp Thr Ser Ile Asn Arg His Asn Asn Phe Arg Thr Phe Leu Gln Ala Leu Met Leu Leu Phe Arg Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys Leu Ser Asn Gln Ala Cys Asp Glu Gln Ala Asn Ala Thr Glu Cys Gly Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu 

Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys Gly Arg

- Ile Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His Met Ser Pro 1745 1750 1755
- Pro Leu Gly Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr Lys 1760 1765 1770
- Arg Leu Val Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr 1775 1780 1785
- Val His Phe Thr Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu 1790 1795 1800
- Glu Ile Lys Leu Ala Pro Ala Gly Thr Lys Gln His Gln Cys Asp 1805 1810 1815
- Ala Glu Leu Arg Lys Glu Ile Ser Val Val Trp Ala Asn Leu Pro 1820 1830
- Gln Lys Thr Leu Asp Leu Leu Val Pro Pro His Lys Pro Asp Glu 1835 1840 1845
- Met Thr Val Gly Lys Val Tyr Ala Ala Leu Met Ile Phe Asp Phe 1850 1855 1860
- Tyr Lys Gln Asn Lys Thr Thr Arg Asp Gln Met Gln Gln Ala Pro 1865 1870 1875
- Gly Gly Leu Ser Gln Met Gly Pro Val Ser Leu Phe His Pro Leu 1880 1885 1890
- Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Val Leu Arg Gly Ala 1895 1900 1905
- Arg Val Phe Leu Arg Gln Lys Ser Ser Thr Ser Leu Ser Asn Gly 1910 1915 1920
- Gly Ala Ile Gln Asn Gln Glu Ser Gly Ile Lys Glu Ser Val Ser 1925 1930 1935
- Trp Gly Thr Gln Arg Thr Gln Asp Ala Pro His Glu Ala Arg Pro 1940 1945 1950

- Pro Leu Glu Arg Gly His Ser Thr Glu Ile Pro Val Gly Arg Ser 1955 1960 1965
- Gly Ala Leu Ala Val Asp Val Gln Met Gln Ser Ile Thr Arg Arg 1970 1975 1980
- Gly Pro Asp Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg 1985 1990 1995
- Ala Ala Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Val Thr 2000 2005 2010
- Asp Ala Ser Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Gln Arg 2015 2020 2025
- Pro Arg Gly Thr His Leu Cys Ser Thr Thr Pro Asp Arg Pro Pro 2030 2035 2040
- Pro Ser Gln Ala Ser Ser His His His His Arg Cys His Arg 2045 2050 2055
- Arg Arg Asp Arg Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu 2060 2065 2070
- Ser Ala Asp Met Asp Gly Ala Pro Ser Ser Ala Val Gly Pro Gly 2075 2080 2085
- Leu Pro Pro Gly Glu Gly Pro Thr Gly Cys Arg Arg Glu Arg Glu 2090 2095 2100
- Arg Arg Gln Glu Arg Gly Arg Ser Gln Glu Arg Arg Gln Pro Ser 2105 2110 2115
- Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr Ser Cys Asp Arg Phe 2120 2125 2130
- Gly Gly Arg Glu Pro Pro Lys Pro Lys Pro Ser Leu Ser Ser His 2135 2140 2145
- Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro His Pro Gln 2150 2155 2160

Ala Gly Ser Ala Val Gly Phe Pro Asn Thr Thr Pro Cys Cys Arg 2165 2170 Glu Thr Pro Ser Ala Ser Pro Trp Pro Leu Ala Leu Glu Leu Ala 2180 2185 Leu Thr Leu Thr Trp Gly Ser Val Trp Thr Val Arg Pro Leu Ser 2195 2200 2205 Thr Pro Cys Leu Arg Thr Arg Ser Leu Ser Arg Arg Leu Trp Pro 2210 2215 2220 Pro Thr Arg Ala Ala Pro Pro Gly Leu Pro Thr Cys Pro Pro 2230 <210> 35 <211> 6176 <212> DNA <213> Homo sapiens <220> <221> exon <222> (1)..(17) <223> Exon 41, continues 5' to nt 1 <220> <221> exon <222> (4510)..(4606) <223> Exon 42 <220> <221> exon <222> (6117)..(6176) <223> Exon 43, continues 3' to nt 6176 <400> 35 tac gac ccg gct gcg tg gtaagtgagc cgtggtgctc tgtggtcctt 47 Tyr Asp Pro Ala Ala Cys gggggtggtc catgcccata gatcttggga tgggcctggt ggcttcagac atgtttcaag 107 tgagcacccc ctcaggtttt ggtttggggc ctagggatct gtcctgtttc gtgtagagca 167 gcagtgatgt cccctccgag cagcacctcc cttttccatc tcgtgtgcct tcgcgcagtg 227 gcccggactg cataaagtga taagggtctt ttgagacctt gagttcagtt ccttgggagg 287 taccaggagt catggaaggt tttcaagggg cttgtgagac tcaggtcttc tgctgaaatt 347 tactgtcttc tttcaaggcc tgtcatgtga gggctcatgg agggcttgtg ggagggtgtg 407

tggagacage acagecatag egteettgtg agactegatg tagggtttta ggacattete 467 cctaaggggc tcaggagtct ccctctgctc tttcttttgg ggacaggggc caacatgcac 527 ctctgtgtcc ccagcccagg ctgcatatag tgtgggtgct tggtgaactt ttttgacaga 587 tgagctacca ggaaggctgt agtgtgttta agacccttgt ctatcatctg tcatcgtggg 647 aaaggtttac ttgtttcaaa ggcgaggggt ggtgcctttc cagacagagg aggagtcgca 707 tgaagggcat ggtttgggtc agaaggtgaa gctgtttgag atcaggatcc atgggggtcc 767 cgtacagcat ggcatctcag cagggctgag agctcattcg ggaaggaaga caggtcaggc 827 agcatggctt ctccatcttt ctgtttgtcc attccttcca tcactcgttc ctcatttact 887 tgctcgctca ctcactcaag caatgtctac tgtgggctgc tccctgcaag gccactaggc 947 ctgctagcac ctcactcaca gcgtgtgccc gggaaatgat cgtggatgag cagctctcag 1007 tgagtgcgta gacatccacc actccccgtg ccctaggggg atgagggaac ygtgggtgcc 1067 accgaggeet geettetete tetgagetee ttageetgee eaggeeeet egeacteete 1127 tgcccccagt gcccttactg ggggagggcc ggcaggacgt ggtcagaccc ctcatggtga 1187 ggtcctactg cccacagage teetgggtte tgcctggttt tgcttaaact gcagetcage 1247 agagggagee etecacagae acaaaaggee tgggageeeg actagaette eecaceetee 1307 tttccactca gcctcccsgg agcaccccca ccttcctagt gaggatccag agggtgctgc 1367 acgggccctt cccagtctgg acaaggtcca gagccactgg cagcatcggg ggccctggga 1427 tggggcctct cctgtccagg ctcttccagg cacacagcat ctctgtgggt gtttttcatt 1487 tgagtcggaa atatacagtt tttacgcagt gcaaaacaga cgtagagatt gtctgtgttt 1547 cacagtgtat atgacaattt ccgtggcttg cgagaaccct gtcatgtccg atgcagcaat 1607 atgtttgccc tgtgatactt acttcatgct gggagggaca ggctcaggca cattcacaca 1667 ggacctgtgg tcaagtaagc ccagtcgacc tgggggagac tgaccacatt gaagacctag 1727 accttgttag gcaaagggca ggagagactc cagccgaggc aggatttagg tgaattctag 1787 gcaggtcacc ctgaatgtgt ggtttagaga gtcgctgctt tttctgggtg tatttttct 1847 rgaggagact gtgttgcaga aaacagagat actttatagc atgtagttag tactraaggg 1907 caaaggcaga tcaacaaagg cctgtgagag gatgacaaca ttccccaatc tttatgagct 1967 acgeceggge ettecacagt cageegeegg caagageaga ggeaggaaga aegggaagtg 2027 2087

ctcagagcag gggtccttag aagacatagg cagtcgggga tgtgagaagt gcttactgaa 2147 gagtaaacaa cagaatacag gcctcacccc agagatagca cgcacaaaat aacacacaaa 2207 acctgcgggc cttgagatag cacagacgtg acctgtgagc aacacatgat tctgccccca 2267 cgagcagcct gggctcaccc tacgtagtcc aagatggcca ggacaggccg aggggcccag 2327 2387 tgageetttt tggtageete tgtgacaaac geagtgatgt agteatttee caggggaeea 2447 ctctatggct cctcgttctt tgggagaaac tttgaaacaa caacaaaatt ttatgagatt 2507 teetgagttg tgtgtgeaca tgtgtggtge acaaacagga catgtgtgca tgtacatgge 2567 tgtgttgtgt gcgtgtttgc acagtgtgac cacatgtgca ctctgtgcat gtgtacatgt 2627 gtgttgtgca cgcatgcaca cagatgtcta catgtgtgtt ttgtgcacat qtqcatqcac 2687 aaggcgcgca cgtgtgtgca tgttgacgtt tgtgtgtgca catatgattg tatatatgtg 2747 tgcctgtgtg tatatgtgtg caagtgaact gaatgtgcgt gcatgcctgt gtggttgtat 2807 gtggatgtgt gtatgaatga catgtgcacc tgtgtgtgca tatgcccatg tgtgcatgtg 2867 tgtgcctgtg gatgtgcaca tgcccgtgtg tgcctgtggt gtgcacgtgt ctgtggtgtg 2927 cacqtgcctg tgtgtgcatg tgcctttgtg tgtgtgccac cgtggtgcgt atgtgcctgt 2987 gtgtgcctgt gggtgtgcac gtgcctgtgg tgtactcgtg cctgtggtgt gcatgtgcct 3047 gtggtgtgca tgtgcctgtg gtgcgcatgt gcctgtgtgg gtgtgcatgt gtgtgcctgt 3107 gggtgtgcac atgtctgtgg tgtacacgtg cctgtggtgc gcatgtgcct gtgtcggtgg 3167 attetgaagg cacgaaactg tteaccetgg tgetgetttt eccageette etattgttgt 3227 tgcttcaacc ctgtggtgtg aagcctggtg acttgagagg agaattgcag tcacacttct 3287 tgcagcaatc tggctgctca tttctgtgtc cacccatctg tctaataaca tctcctgggc 3347 atctgccatg tgctggggca ggggcccaaa ggtgaaacag gcacatgctg tcagaacact 3407 cagccaggct tggcagtgag accacatggt gactgtagct caaggccagg gcatttctqa 3467 ggaaggagag gccatgtctg gcctgaagaa tgagaggagg tgccatttga gctgaactta 3527 aaggaatagg aaggattgag cactagagaa ataggatgga ttgagctggg gatgggaggt 3587 ggggcagagg gaagagggag gccaggtggc ttctgtcagg ctggagagga gagtgcattg 3647 cagaggggct gcaaggaagg ctgagaccgt ttgagaaagg cattgaaatt ggttgaacaa 3707 gtegetttga ttttgegttg tagacagtgg geteagteet tgeategagg agtgatggtg 3767 tcaggcccac actttccacc tgggcttgat gggcagggac agctggggag ggtgggggt 3827

ctcacgtgtg tccagcccc cgtggtggtc ccttggctgc cacaacctta gcaaacatct	3887
ttggtgaggc cgaggtgggg cttcctgttt cttccctttt acagaaggaa attcaacatg	3947
atttggaget gattggaceg agggggetae actgtaggag agaggageat tggtggtgat	4007
ccaaggatge cageetacga tgagaggagg teettagetg gaecaggget teeaggggag	4067
gacacagagt gttcagggag acagggtcag tcaggaggca aatgggtatg ggcctctgag	4127
gttgctgtga tagagaagca gacctgcctt gctgtggtgg tgagaaagcc aggcgtcaga	4187
gggtttggct gacctgggag gcccagggag catcgtcact cagggagtga gaggaagagc	4247
tgaccgagag ggaggggcag ggcagtgttc tgtgtggagc ggctcctgca cacatcgggc	4307
teeggaaatt tegetggggt tgeeacaeet eeeggggage teeaageeee agtaceeegg	4367
cgaggtgcca cctgtccagt gcacccctca cctccgccgc cacgcctgcg cgaggtcggt	4427
cttcccatca caggctgggc gggcgggagg gtcgggctcc atcctgcctt tgttgccata	4487
acctgtgaca tttcctttcc ag t tgc cgg att cat tat aag gat atg tac Cys Arg Ile His Tyr Lys Asp Met Tyr 10 15	4537
agt ttg ttg cgt tgt att gcg cca ccc gtt ggc tta ggg aag aac tgc Ser Leu Leu Arg Cys Ile Ala Pro Pro Val Gly Leu Gly Lys Asn Cys	4585
20 25 30	
	4636
20 25 30  cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat  Pro Arg Arg Leu Ala Tyr Lys	4636 4696
20 25 30  cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat  Pro Arg Arg Leu Ala Tyr Lys  35	
20 25 30  cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat  Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc	4696
20 25 30  cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat  Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc  ctcacctttc tttcctcctg tcctgccccg cactccagcc tcctggacgc tttggttccc	4696 4756
cct cgt agg ttg gcc tac aag gtttgcaact tcagagcetc catccaacat Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc ctcacctttc tttcctcctg tcctgcccg cactccagcc tcctggacgc tttggttccc tgtctaatct cccctgaccc ttctagctgg ctggggacag gcagcctttt gagagtgcac	4696 4756 4816
cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc ctcacctttc tttcctcctg tcctgccccg cactccagcc tcctggacgc tttggttccc tgtctaatct cccctgaccc ttctagctgg ctggggacag gcagccttt gagagtgcac agaaagcttt ctggttggga tcagctagct gggtgggaga tgggggctag gtccctcctg	4696 4756 4816 4876
cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc ctcacctttc tttcctcctg tcctgcccg cactccagcc tcctggacgc tttggttccc tgtctaatct cccctgaccc ttctagctgg ctggggacag gcagccttt gagagtgcac agaaagcttt ctggttggga tcagctagct gggtgggaga tgggggctag gtccctcctg catcaaggaa gcagccgact tgcccgttt ttcctccgag ctctcgagag tgagctgatg	4696 4756 4816 4876 4936
cct cgt agg ttg gcc tac aag gtttgcaact tcagagcctc catccaacat Pro Arg Arg Leu Ala Tyr Lys 35  acttccctgc ccgaaacgca cacagcacac actccctgcc tggtttggga atgacgacgc ctcacctttc tttcctcctg tcctgcccg cactccagcc tcctggacgc tttggttccc tgtctaatct cccctgaccc ttctagctgg ctggggacag gcagccttt gagagtgcac agaaagcttt ctggttggga tcagctagct gggtgggaga tgggggctag gtccctcctg catcaaggaa gcagccgact tgcccggttt ttcctccgag ctctcgagag tgagctgatg gtcagccaa aggccaa aggccgagg ccccagagga agggagcacg ggcacaggcc tcccaggagc	4696 4756 4816 4876 4936
cet egt agg ttg gee tae aag gtttgeaact teagageete eatecaacat Pro Arg Arg Leu Ala Tyr Lys 35  actteeetge eegaaacgea cacageacae acteeetgee tggtttggga atgacgaege etcacettte ttteeteetg teetgeeeg eacteeagee teetggaege tttggtteee tgtetaatet eeeetgaeee ttetagetgg etggggaeag geageettt gagagtgeae agaaagett etggttggga teagetaget gggtgggaga tgggggetag gteeeteetg eateaaggaa geageegaet tgeeeggtt tteeteegag etetegagag tgagetgatg gteageegaegeegaegeegaeggeegaegggaegg	4696 4756 4816 4876 4936 4996 5056
cet egt agg ttg gee tac aag gtttgeaact teagageete eatecaacat Pro Arg Arg Leu Ala Tyr Lys 35 actteetge eegaaacgea cacageacae acteeetgee tggtttgga atgaegaege etcacettte ttteeteetg teetgeeeeg eacteeagee teetggaege tttggtteee tgtetaatet eccetgaece ttetagetgg etggggaeag geageettt gagagtgeae agaaagettt etggttgga teagetaget gggtggaga tgggggetag gteeeteetg eateaaggaa geageegaet tgeeeggtt tteeteegag etetegagag tgagetgatg gteaggeeaa aggeegaea aggeegaeg ecceagaga agggageaeg ggeaeaggee teecaggag ageegteete tgggttggaa gageecagta acceagtgge tetecagtee etgggggagg aaagggeaga ggeagaggg gggeeaaaga eaageegtee tgetttgeet etgggggagg	4696 4756 4816 4876 4936 4996 5056 5116

gcctcgtggc ctctctactt gggaagactc tgtcatcgga gcctttaggt tcctagatcc	5356
tgcaacctgc ccctttgctg tcagagctgg tgcaaggtgg ccccaggcca tgtcctgcca	5416
getectecce cetecegeag teteceattt cecaeceage eetgageece ageageagee	5476
ctgccctggg ctggcctcgg gcgtctctgg gcttgaagca ggacttgccc caaggttagg	5536
ttttggctgt acagettcag ggetceetgt ttagttttet ggtttgttte agatettete	5596
cacettttta geetetagee teaettggaa geaaatetga eegtgaggaa agetgatgte	5656
tetgeegtgg tteteageea etgeeteate tgeeacetga ecetgeetge eagetettet	5716
ggcagggtgg accagctett ttetgattgg catteactgt gatttttaaa agagtaaaat	5776
cagaaattat caacccaaag ccgccccgat gcagcccttc ctcacccctc catgttcatt	5836
ageteagaeg ceacecege ceactgeece gegtgggget ggagtgagga ggtgaggete	5896
gggggtgggg ggccaggagg gggtcaagcc aagcagagaa atccctcatt ttatttattt	5956
tgattttggg ctttgaatcc gacactttga ttaactggct ctgttttctt gtctgcttcc	6016
tecetgeece tacceegete eceteceege teceeteetg etgeegetee teetgtggae	6076
caccccactg tgccctggcc tcgctgggac gggtttccag tgg gcg cat cag tta Trp Ala His Gln Leu 40	6131
caa tga cat gtt tga gat gct gaa aca cat gtc ccc gcc tct ggg Gln His Val Asp Ala Glu Thr His Val Pro Ala Ser Gly 45 50 55	6176

<210> 36

<211> 38 <212> PRT

<213> Homo sapiens

Tyr Asp Pro Ala Ala Cys Cys Arg Ile His Tyr Lys Asp Met Tyr Ser 1 5

Leu Leu Arg Cys Ile Ala Pro Pro Val Gly Leu Gly Lys Asn Cys Pro 20 . 25 30

Arg Arg Leu Ala Tyr Lys 35

<210> 37 <211> 6984 <212> DNA

## <213> Mus musculus

<220> <221> CDS <222> (1)(6984)	
<pre>&lt;400&gt; 37 atg gtc cgc ttc ggg gac gag cta ggc ggc cgc tat ggg gg Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gl 1 5 10</pre>	
ggc ggg gag cgg gct cgg ggc ggc ggc ggc ggc ggg gcg gg Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gl 20 25 30	y Gly Pro
ggc cag ggg ggt ctg ccg ccg ggc cag cgg gtc ctg tac aa Gly Gln Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Ly 35 40 45	
att gcg cag cgc gca cgg act atg gcc ctg tac aac ccc at Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Il 50 55 60	_
aag cag aac tgc ttc acc gtc aac cgc tcg ctc ttc gtc tt Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Ph 65 70 75	5 5 5
gac aac gtc gtc cgc aaa tac gct aag cgc atc acc gaa tg Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Tr 85 90	
ttc gaa tac atg atc ctg gcc acc atc atc gcc aac tgt at Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Il 100 105 11	e Val Leu
gcc ctg gag cag cac ctc cct gat ggg gac aag act ccc at Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Me 115 120 125	
cga cta gat gac acg gag cct tac ttc atc ggg atc ttt tg Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cy 130 135 140	
gcg ggc atc aag atc ata gcc ctg ggc ttt gtt ttc cac aa Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Ly 145	
tac ctt cgg aac ggc tgg aat gtc atg gac ttc gtg gtg gt Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Va 165 170	
ggg att ctc gcc aca gct gga act gac ttt gac ctg cgc ac Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Th 180 185 19	r Leu Arg
gct gtg cgt gtg ctt agg ccc ctg aag ctg gtg tct gga at	t cca agc 624

Ala	Val	Arg 195	Val	Leu	Arg	Pro	Leu 200	Lys	Leu	Val	Ser	Gly 205	Ile	Pro	Ser	
							atc Ile									672
_	Ile		_	_			ttt Phe	_			_		_			720
							aaa Lys									768
_		_					ggt Gly	_			_			_		816
							gac Asp 280									864
							aat Asn									912
							acc Thr									960
							ggc Gly									1008
							tcc Ser			_			_		_	1056
							gcc Ala 360									1104
cgc Arg	cgc Arg 370	gcc Ala	ttc Phe	ctg Leu	aag Lys	ctc Leu 375	cgc Arg	agg Arg	cag Gln	cag Gln	cag Gln 380	att Ile	gag Glu	cga Arg	gag Glu	1152
							atc Ile									1200
							gaa Glu									1248
	_	_	_		-	_	agc Ser	_		_				_	_	1296

.

420 425 430

gag ggg gag ga Glu Gly Glu As 435		_	Cys Ala Val		
gct cgt gcc ag Ala Arg Ala Se 450		Gly Lys T			
cgg aga aag ga Arg Arg Lys Gl 465					
gca cag agc tt Ala Gln Ser Ph		. Val Leu C			n Thr
ctg tgt gtg gc Leu Cys Val Al 50	a Met Val His		-		
gca ctg tac tt Ala Leu Tyr Ph 515			eu Gly Leu		
atg tcc ctg aa Met Ser Leu Ly 530		Leu Gly P		-	-
tcc ttc aac tg Ser Phe Asn Cy 545					
gta gtc tgg gc Val Val Trp Al		Pro Gly T			r Val
ctg cgg gct ct Leu Arg Ala Le 58	Arg Leu Lei		_	-	
aac tct ctg ag Asn Ser Leu Ard 595			eu Leu Asn	_	•
atc atc agc ct Ile Ile Ser Le 610		Leu Phe L			
ctg ttg ggg atc Leu Leu Gly Me 625					
act cca acc acc Thr Pro Thr Th		Thr Phe P			r Val

	_		_			gag Glu	-			_	_	_				:	2016
		_				gtc Val	_			_						:	2064
		-	_		_	ttt Phe 695					_	_				:	2112
_	_		_		_	aac Asn		_		_	_		_		_	:	2160
						gaa Glu										:	2208
_	_		_	-	-	gaa Glu	_	_		_		_	_			:	2256
						cag Gln										:	2304
	_		_	_	_	cta Leu 775			_		_	_		~	_	:	2352
						atg Met										:	2400
_	_	_				cca Pro	_	. –	_			_		_		:	2448
						cga Arg										:	2496
_		-		_		acg Thr		-		_	_		_			:	2544
_						cgt Arg 855	_		_	_			_		_	:	2592
						gac Asp										:	2640

cgg gag gaa cgt gcg cgc cct cgt cga agt cac agc aag gag act cca Arg Glu Glu Arg Ala Arg Pro Arg Arg Ser His Ser Lys Glu Thr Pro 885 890 895	2688
ggg gct gac acg caa gtg cgc tgt gag cgc agt agg cgt cac cac cgg Gly Ala Asp Thr Gln Val Arg Cys Glu Arg Ser Arg Arg His His Arg 900 905 910	2736
cgc ggc tcc ccg gag gag gcc act gaa cgg gag cct cgg cgc cac cgt Arg Gly Ser Pro Glu Glu Ala Thr Glu Arg Glu Pro Arg Arg His Arg 915 920 925	2784
gcc cac cgg cat gca cag gac tca agc aag gag ggc acg gcg ccg gtg Ala His Arg His Ala Gln Asp Ser Ser Lys Glu Gly Thr Ala Pro Val 930 935 940	2832
ctt gta ccc aag ggt gag cga cga gca aga cac cga ggc cca cgc acg Leu Val Pro Lys Gly Glu Arg Arg Ala Arg His Arg Gly Pro Arg Thr 945 950 955 960	2880
ggt cca cgt gag gca gag aac aac gag gag ccc aca cgc agg cac cgt Gly Pro Arg Glu Ala Glu Asn Asn Glu Glu Pro Thr Arg Arg His Arg 965 970 975	2928
gca agg cat aag gtg cca ccc aca ctg cag ccc cca gag agg gag gct Ala Arg His Lys Val Pro Pro Thr Leu Gln Pro Pro Glu Arg Glu Ala 980 985 990	2976
gca gag aag gag agc aac gcg gtg gaa ggg gat aag gaa acc cga aat Ala Glu Lys Glu Ser Asn Ala Val Glu Gly Asp Lys Glu Thr Arg Asn 995 1000 1005	3024
cac cag ccc aag gaa cct cac tgt gac ctg gag gcc att gca gtt His Gln Pro Lys Glu Pro His Cys Asp Leu Glu Ala Ile Ala Val 1010 1015 1020	3069
aca ggt gtg ggc cct ctg cac atg ctg ccc agc acc tgt ctc cag Thr Gly Val Gly Pro Leu His Met Leu Pro Ser Thr Cys Leu Gln 1025 1030 1035	3114
aaa gtg gac gag caa cca gag gat gca gac aac cag cgt aat gtc Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val 1040 1045 1050	3159
Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val	3159 3204
Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val 1040 1045 1050  acc cgg atg ggc agt cag ccc tca gat ccc agc acc act gtg cat Thr Arg Met Gly Ser Gln Pro Ser Asp Pro Ser Thr Thr Val His	
Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln 1040  acc cgg atg ggc agt cag ccc tca gat ccc agc acc act gtg cat Thr Arg Met Gly Ser Gln Pro 1060  gtc cca gtg aca ctg aca ggc cct cct ggg gag acc cct gta gtt Val Pro Val Thr Leu Thr Gly Pro Pro Gly Glu Thr Pro Val Val	3204

Glu	Ala 1100	Glu	Ala	Asp	Asp	Val 1105	Leu	Arg	Arg	Gly	Pro 1110	Arg	Pro	Ile		
_	ccc Pro 1115										acc Thr 1125				3	384
_	cgc Arg 1130										tac Tyr 1140				3	429
_	att Ile 1145					_	_	_	_		gcc Ala 1155	_	_	_	3	474
											aac Asn 1170				3	519
	atg Met 1175	-						-				_	_	_	3	564
	aag Lys 1190	_		_	-		_	_	_		cct Pro 1200		_		3	609
	cgg Arg 1205										gtt Val 1215				3	654
<b>-</b>	ctg Leu 1220										gga Gly 1230				3	699
	aaa Lys 1235														3	744
~				_			_		_		aag Lys 1260		-	-	3	789
_	ttt Phe 1265										gtc Val 1275				3	834
_	att Ile 1280	_		_			_				_	_			3	879
_	cag Gln 1295					_				_	act Thr 1305	_	_		3	924
_	gag Glu										ttg Leu				3	969

1310				1315			1320	
 J	 _	_	_	cag Gln		 		1
1325				1330			1335	

						cag Gln 1330									4014
	cac His 1340		_		_	ctc Leu 1345	Trp	_	_	_	_	_			4059
	tcc Ser 1355					tgg Trp 1360									4104
						cag Gln 1375									4149
						gtg Val 1390									4194
						gtg Val 1405									4239
	cag Gln 1415					atg Met 1420					tta Leu 1425				4284
						ttt Phe 1435									4329
	tac Tyr 1445					aaa Lys 1450					tat Tyr 1455				4374
						ccc Pro 1465									4419
		Leu		Thr	Val	gtg Val 1480	Leu	Met	Met	Lys		Tyr			4464
						atg Met 1495									4509
aca Thr	tcc Ser 1505	atg Met	ttc Phe	tcg Ser	atg Met	gag Glu 1510	tgc Cys	ata Ile	ctg Leu	aag Lys	atc Ile 1515	atc Ile	gcc Ala	ttt Phe	4554
						aga Arg 1525									4599

_	acg Thr 1535	_	_		_			_		_				4644
	aac Asn 1550													4689
_	cgg Arg 1565	-		_	_		_	_				_		4734
	ttg Leu 1580					_			_	 _				4779
	ctc Leu 1595													4824
	cag Gln 1610													4869
	cga Arg 1625													4914
	ttc Phe 1640										Ile			4959
	tgt Cys 1655	_				_		_		_		_	_	5004
	tgc Cys 1670											ttc Phe	_	5049
	ctc Leu 1685	_			_	_	_			_	_	_		5094
_	gac Asp 1700								_					5139
	cac His 1715													5184
	gct Ala 1730													5229

Leu l			_			cct Pro 1750	_		_		_		_	_	5274
Ala A	_	_	_		-	cgc Arg 1765	_	_	_	_		_	ccc Pro		5319
Ser i			_	_	_	gtg Val 1780			_			_	atg Met	_	5364
Leu :						gag Glu 1795									5409
Lys (						gct Ala 1810							tct Ser		5454
Val '						cag Gln 1825									5499
Pro 1				_		atg Met 1840				_	_		gct Ala	_	5544
Leu l	atg Met 1850			_		tac Tyr 1855		_					aga Arg	-	5589
Gln '				_		gga Gly 1870		_		_	_		ccc Pro		5634
Ser :	_				_	aag Lys 1885	_		_	-	_		cag Gln		5679
Ala '		Leu	Arg	Gly	Ala	cgg Arg 1900	Val	Phe	Leu	Arg		Lys			5724
Thr	tcc Ser 1910	ctc Leu	agc Ser	aat Asn	gly aaa	ggt Gly 1915	gcc Ala	ata Ile	caa Gln	acc Thr	cag Gln 1920	gaa Glu	agt Ser	ggc Gly	5769
Ile I	_		-,-	_		tgg Trp 1930		_	_				_	-	5814
Leu '						cct Pro 1945									5859
atc	cct	gtg	999	cag	tca	gga	aca	ctg	gct	gtg	gat	gtc	cag	atg	5904

Ile		Val	Gly	Gln	Ser	Gly 1960	Thr	Leu	Ala	Val	Asp 1965	Val	Gln	Met	
						gga Gly 1975									5949
	gaa Glu 1985	_			_	gct Ala 1990	~		_		_			_	5994
-		_	_	_		aat Asn 2005	_	_		_	_	_			6039
						cca Pro 2020									6084
						cct Pro 2035									6129
						aga Arg 2050									6174
						gtt Val 2065									6219
						ccc Pro 2080									6264
						gag Glu 2095									6309
						tca Ser 2110									6354
						gag Glu 2125									6399
						cca Pro 2140							_		6444
						tcc Ser 2155									6489
						ccg Pro									6534

	2165					2170			2175				
	cag Gln 2180					cca Pro 2185							6579
_	aat Asn 2195		_		_	cac His 2200	_	 	_	_	ggc Gly		6624
	gcc Ala 2210					cgt Arg 2215					gaa Glu		6669
						gag Glu 2230							6714
						gac Asp 2245				-	ctg Leu	_	6759
						acc Thr 2260					acc Thr		6804
	gag Glu 2270					aac Asn 2275					act Thr		6849
	gtg Val 2285					tcc Ser 2290							6894
	aat Asn 2300					act Thr 2305					gtc Val		6939
						cac His 2320					_	tag	6984
<210 <211 <212 <213	L> 23 2> PF	327 RT	ıscu:	lus									

<400> 38

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Thr Gly 1 5 10 15

- Gly Gln Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45
- Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50 55 60
- Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80
- Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95
- Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110
- Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125
- Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140
- Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser 145 150 155 160
- Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr
  165 170 175
- Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 185 190
- Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205
- Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 215 220
- Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile 225 230 235 240
- Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 255

- Ser Thr Asp Thr Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Asp Pro 260 265 270
- Pro Ala Arg Gln Cys Asp Gly Asp Thr Glu Cys Arg Glu Tyr Trp Pro 275 280 285
- Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 290 295 300
- Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 305 310 315 320
- Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 325 330 335
- Ile Pro Leu Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 350
- Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365
- Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 380
- Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400
- Ala Glu Glu Asp Lys Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu 405 410 415
- Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu 420 425 430
- Glu Gly Glu Asp Arg Phe Val Asp Leu Cys Ala Val Gly Ser Pro Phe 435 440 445
- Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe 450 455 460
- Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys 465 470 475 480
- Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr

185	490	495

Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Gln Arg Leu Thr Thr

500 505 Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser 530 535 Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe Glu 545 550 555 Val Val Trp Ala Ala Ile Lys Pro Gly Thr Ser Phe Gly Ile Ser Val Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp 585 Asn Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 600 Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala 615 Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val 645 650 Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly 670

Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr 675 680 685

Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe

Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys

700

715

695

710

- Asp Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln
  725 730 735
- Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750
- Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp 755 760 765
- Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys
  770 780
- Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Tyr Ala 785 790 795 800
- Ser Thr Arg His Val Arg Pro Asp Met Lys Thr His Met Asp Arg Pro 805 810 815
- Leu Val Val Glu Pro Gly Arg Asp Gly Leu Arg Gly Pro Val Gly Ser 820 825 830
- Lys Ser Lys Pro Glu Gly Thr Glu Ala Thr Glu Ser Ala Asp Leu Pro 835 840 845
- Arg Arg His His Arg His Arg Asp Arg Asp Lys Thr Ser Ala Thr Ala 850 855 860
- Pro Ala Gly Gly Glu Gln Asp Arg Thr Glu Ser Thr Glu Thr Gly Ala 865 870 875 880
- Arg Glu Glu Arg Ala Arg Pro Arg Arg Ser His Ser Lys Glu Thr Pro 885 890 895
- Gly Ala Asp Thr Gln Val Arg Cys Glu Arg Ser Arg Arg His His Arg 900 905 910
- Arg Gly Ser Pro Glu Glu Ala Thr Glu Arg Glu Pro Arg Arg His Arg 915 920 925
- Ala His Arg His Ala Gln Asp Ser Ser Lys Glu Gly Thr Ala Pro Val 930 935 940

- Leu Val Pro Lys Gly Glu Arg Arg Ala Arg His Arg Gly Pro Arg Thr 945 950 955 960
- Gly Pro Arg Glu Ala Glu Asn Asn Glu Glu Pro Thr Arg Arg His Arg 965 970 975
- Ala Arg His Lys Val Pro Pro Thr Leu Gln Pro Pro Glu Arg Glu Ala 980 985 990
- Ala Glu Lys Glu Ser Asn Ala Val Glu Gly Asp Lys Glu Thr Arg Asn 995 1000 1005
- His Gln Pro Lys Glu Pro His Cys Asp Leu Glu Ala Ile Ala Val 1010 1015 1020
- Thr Gly Val Gly Pro Leu His Met Leu Pro Ser Thr Cys Leu Gln 1025 1030 1035
- Lys Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val 1040 1045 1050
- Thr Arg Met Gly Ser Gln Pro Ser Asp Pro Ser Thr Thr Val His 1055 1060 1065
- Val Pro Val Thr Leu Thr Gly Pro Pro Gly Glu Thr Pro Val Val 1070 1075 1080
- Pro Ser Gly Asn Met Asn Leu Glu Gly Gln Ala Glu Gly Lys Lys 1085 1090 1095
- Glu Ala Glu Ala Asp Asp Val Leu Arg Arg Gly Pro Arg Pro Ile 1100 1105 1110
- Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Leu 1115 1120 1125
- Arg Arg Phe Cys His Tyr Ile Val Thr Met Arg Tyr Phe Glu Met 1130 1140
- Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile Ala Leu Ala Ala 1145 1150 1155

- Glu Asp Pro Val Arg Thr Asp Ser Phe Arg Asn Asn Ala Leu Lys 1160 1165 1170
- Tyr Met Asp Tyr Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val 1175 1180 1185
- Ile Lys Met Ile Asp Leu Gly Leu Leu His Pro Gly Ala Tyr 1190 1195 1200
- Phe Arg Asp Leu Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly 1205 1210
- Ala Leu Val Ala Phe Ala Phe Ser Ser Phe Met Gly Gly Ser Lys 1220 1225 1230
- Gly Lys Asp Ile Asn Thr Ile Lys Ser Leu Arg Val Leu Arg Val 1235 1240 1245
- Leu Arg Pro Leu Lys Thr Ile Lys Arg Leu Pro Lys Leu Lys Ala 1250 1260
- Val Phe Asp Cys Val Val Asn Ser Leu Lys Asn Val Leu Asn Ile 1265 1270 1275
- Leu Ile Val Tyr Met Leu Phe Met Phe Ile Phe Ala Val Ile Ala 1280 1285 1290
- Val Gln Leu Phe Lys Gly Lys Phe Phe Tyr Cys Thr Asp Glu Ser 1295 1300 1305
- Lys Glu Leu Glu Arg Asp Cys Arg Gly Gln Tyr Leu Asp Tyr Glu 1310 1315 1320
- Lys Glu Glu Val Glu Ala Gln Pro Arg Gln Trp Lys Lys Tyr Asp 1325 1330 1335
- Phe His Tyr Asp Asn Val Leu Trp Ala Leu Leu Thr Leu Phe Thr 1340 1350
- Val Ser Thr Gly Glu Gly Trp Pro Met Val Leu Lys His Ser Val 1355 1360 1365
- Asp Ala Thr Tyr Glu Glu Gln Gly Pro Ser Pro Gly Phe Arg Met

1370 1375 . 1380

Glu	Leu 1385		Ile	Phe	Tyr	Val 1390		Tyr	Phe	Val	Val 1395	Phe	Pro	Phe
Phe	Phe 1400	Val	Asn	Ile	Phe	Val 1405	Ala	Leu	Ile	Ile	Ile 1410	Thr	Phe	Gln
Glu	Gln 1415	Gly	Asp	Lys	Val	Met 1420		Glu	Cys	Ser	Leu 1425	Glu	Lys	Asn
Glu	Arg 1430	Ala	Cys	Ile	Asp	Phe 1435	Ala	Ile	Ser	Ala	Lys 1440	Pro	Leu	Thr
Arg	Tyr 1445	Met	Pro	Gln	Asn	Lys 1450		Ser	Phe	Gln	Tyr 1455	Lys	Thr	Trp
Thr	Phe 1460	Val	Val	Ser	Pro	Pro 1465	Phe	Glu	Tyr	Phe	Ile 1470	Met	Ala	Met
Ile	Ala 1475	Leu	Asn	Thr	Val	Val 1480	Leu	Met	Met	Lys	Phe 1485	Tyr	Asp	Ala
Pro	Tyr 1490	Glu	Tyr	Glu	Leu	Met 1495	Leu	Lys	Cys	Leu	Asn 1500	Ile	Val	Phe
Thr	Ser 1505	Met	Phe	Ser	Met	Glu 1510	Cys	Ile	Leu	Lys	Ile 1515	Ile	Ala	Phe
Gly	Val 1520	Leu	Asn	Tyr	Phe	Arg 1525	Asp	Ala	Trp	Asn	Val 1530	Phe	Asp	Phe
Val	Thr 1535	Val	Leu	Gly	Ser	Ile 1540		Asp	Ile	Leu	Val 1545	Thr	Glu	Ile
Ala	Asn 1550	Asn	Phe	Ile	Asn	Leu 1555	Ser	Phe	Leu	Arg	Leu 1560	Phe	Arg	Ala
Äla	Arg 1565	Leu	Ile	Lys	Leu	Leu 1570	Arg	Gln	Gly	Tyr	Thr 1575	Ile	Arg	Ile
Leu	Leu 1580	Trp	Thr	Phe	Val	Gln 1585	Ser	Phe	Lys	Ala	Leu 1590	Pro	Tyr	Val

- Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile Gly 1595 1600 1605
- Met Gln Val Phe Gly Asn Ile Ala Leu Asp Asp Asp Thr Ser Ile 1610 1615 1620
- Asn Arg His Asn Asn Phe Arg Thr Phe Leu Gln Ala Leu Met Leu 1625 1630 1635
- Leu Phe Arg Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu 1640 1650
- Ser Cys Leu Gly Asn Arg Ala Cys Asp Pro His Ala Asn Ala Ser 1655 1660 1665
- Glu Cys Gly Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile 1670 1680
- Phe Leu Cys Ser Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile 1685 1690 1695
- Met Asp Asn Phe Glu Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly 1700 1705 1710
- Pro His His Leu Asp Glu Phe Ile Arg Val Trp Ala Glu Tyr Asp 1715 1720 1725
- Pro Ala Ala Cys Gly Arg Ile Ser Tyr Asn Asp Met Phe Glu Met 1730 1740
- Leu Lys His Met Ser Pro Pro Leu Gly Leu Gly Lys Lys Cys Pro 1745 1750 1755
- Ala Arg Val Ala Tyr Lys Arg Leu Val Arg Met Asn Met Pro Ile 1760 1765 1770
- Ser Asn Glu Asp Met Thr Val His Phe Thr Ser Thr Leu Met Ala 1775 1780 1785
- Leu Ile Arg Thr Ala Leu Glu Ile Lys Leu Ala Pro Ala Gly Thr 1790 1795 1800

- Lys Gln His Gln Cys Asp Ala Glu Leu Arg Lys Glu Ile Ser Ser 1805 1810 1815
- Val Trp Ala Asn Leu Pro Gln Lys Thr Leu Asp Leu Leu Val Pro 1820 1825 1830
- Pro His Lys Pro Asp Glu Met Thr Val Gly Lys Val Tyr Ala Ala 1835 1840 1845
- Leu Met Ile Phe Asp Phe Tyr Lys Gln Asn Lys Thr Thr Arg Asp 1850 1855 1860
- Gln Thr His Gln Ala Pro Gly Gly Leu Ser Gln Met Gly Pro Val 1865 1870 1875
- Ser Leu Phe His Pro Leu Lys Ala Thr Leu Glu Gln Thr Gln Pro 1880 1885 1890
- Ala Val Leu Arg Gly Ala Arg Val Phe Leu Arg Gln Lys Ser Ala 1895 1900 1905
- Thr Ser Leu Ser Asn Gly Gly Ala Ile Gln Thr Gln Glu Ser Gly 1910 1915 1920
- Ile Lys Glu Ser Leu Ser Trp Gly Thr Gln Arg Thr Gln Asp Ala 1925 1930 1935
- Leu Tyr Glu Ala Arg Ala Pro Leu Glu Arg Gly His Ser Ala Glu 1940 1945 1950
- Ile Pro Val Gly Gln Ser Gly Thr Leu Ala Val Asp Val Gln Met 1955 1960 1965
- Gln Asn Met Thr Leu Arg Gly Pro Asp Gly Glu Pro Gln Pro Gly 1970 1980
- Leu Glu Ser Gln Gly Arg Ala Ala Ser Met Pro Arg Leu Ala Ala 1985 1990 1995
- Glu Thr Gln Pro Ala Pro Asn Ala Ser Pro Met Lys Arg Ser Ile 2000 2005 2010

- Ser Thr Leu Ala Pro Arg Pro His Gly Thr Gln Leu Cys Ser Thr 2015 2020 2025
- Val Leu Asp Arg Pro Pro Pro Ser Gln Ala Ser His His His His 2030 2035 2040
- His Arg Cys His Arg Arg Arg Asp Lys Lys Gln Arg Ser Leu Glu 2045 2050 2055
- Lys Gly Pro Ser Leu Ser Val Asp Pro Glu Gly Ala Pro Ser Thr 2060 2065 2070
- Ala Ala Gly Pro Gly Leu Pro His Gly Glu Gly Ser Thr Ala Cys 2075 2080 2085
- Arg Arg Asp Arg Lys Gln Glu Arg Gly Arg Ser Gln Glu Arg Arg 2090 2095 2100
- Gln Pro Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr Ser Cys 2105 2110 2115
- Asp Arg Phe Gly Ser Arg Glu Pro Pro Gln Leu Met Pro Ser Leu 2120 2125 2130
- Ser Ser His Pro Thr Ser Pro Thr Ala Ala Leu Glu Pro Ala Pro 2135 2140 2145
- His Pro Gln Gly Ser Gly Ser Val Asn Gly Ser Pro Leu Met Ser 2150 2155 2160
- Thr Ser Gly Ala Ser Thr Pro Gly Arg Gly Gly Arg Arg Gln Leu 2165 2170 2175
- Pro Gln Thr Pro Leu Thr Pro Arg Pro Ser Ile Thr Tyr Lys Thr 2180 2185 2190
- Ala Asn Ser Ser Pro Val His Phe Ala Glu Gly Gln Ser Gly Leu 2195 2200 2205
- Pro Ala Phe Ser Pro Gly Arg Leu Ser Arg Gly Leu Ser Glu His 2210 2215 2220
- Asn Ala Leu Leu Gln Lys Glu Pro Leu Ser Gln Pro Leu Ala Pro

2225 2230 2235

Gly	Ser 2240	Arg	Ile	Gly	Ser	Asp 2245		ту	r L	eu	Gly	Gln 2250	_	Leu	Asp	
Ser	Glu 2255		Ser	Ala	His	Thr 2260		ı Pr	o G	lu	Asp	Thr 2265		Thr	Phe	
Glu	Glu 2270	Ala	Val	Ala	Thr	Asn 2275		gl	у А	ırg	Ser	Ser 2280	_	Thr	Ser	
Tyr	Val 2285		Ser	Leu	Thr	Ser 2290		n Se	r H	lis	Pro	Leu 2295	_	Arg	Val .	
Pro	Asn 2300	_	Tyr	His	Cys	Thr 2305		ı Gl	у L	eu	Ser	Thr 2310	-	Val	Arg	
Ala	Arg 2315		Ser	Tyr	His	His 2320		As	p G	ln	Asp	His 2325	-	Cys		
		185 NA	uscu	lus										٠_		
<220 <221 <222	L> C	DS 121)	(6	987)												
<400 ggaa			cgag	gggc	g ag	gtcca	ggc	ago	tcg	ıctg	ac di	gctaç	ıgcta	ggag	geeettg	60
gcg	gccg	cg c	cctc	ggtg	c cg	ggccg	cgg	ago	ccg	ıgga	at go	ctcgo	ggcg	ccc	gggagtc	120
			Phe (												ggc Gly	168
		Glu				ggc g Gly G	ly G									216
	Gln					ccg g Pro G 4	ly c						r Ly			264
					Arg '	act a Thr M 55						sn Pi				312

_	_		_			_	aac Asn	_	_			-		_		360
_		_	-	_			gct Ala	_	_			_		_		408
	_		_		_	_	acc Thr					_		_	_	456
		_	_				gat Asp 120		-	_			_			504
_		_	_	_			tac Tyr						_			552
							ctg Leu									600
							gtc Val	_	_			~ ~	_		_	648
			-		_		act Thr	_		_	_	_		_		696
							ctg Leu 200									744
							atc Ile			-	_	_	_	-	_	792
	_		_	_			ttt Phe				_		-			840
					_		aaa Lys			_	_	_				888
							ggt Gly									936
							gac Asp 280									984

								aat Asn									1032
]	_				_	_		acc Thr	_					_			1080
					_		_	ggc Gly		_				_			1128
								tcc Ser			_			_		ctg ' Leu	1176
								gcc Ala 360									1224
								cgc Arg									1272
:								atc Ile									1320
								gaa Glu	_				_	_		_	1368
	_	_	_	-		_	_	agc Ser	_							_	1416
				_			_	gac Asp 440		_	_	_					1464
								Gly aaa									1512
į								cgg Arg									1560
								gta Val									1608
	_	_		_	_			tat Tyr		_		_					1656
9	gca	ctg	tac	ttt	gca	gag	ttt	gtt	ttc	ctg	ggt	ctc	ttc	ctc	aca	gag	1704

Ala Leu	Tyr Pho	e Ala	Glu	Phe	Val 520	Phe	Leu	Gly	Leu	Phe 525	Leu	Thr	Glu		
atg tcc Met Ser 530	_	_						_	_					1752	
tcc ttc Ser Phe 545														1800	
gta gtc Val Val														1848	
ctg cgg Leu Arg		u Arg												1896	
aac tct Asn Ser														1944	
atc atc Ile Ile 610														1992	
ctg ttg Leu Leu 625														2040	
act cca Thr Pro				Asp										2088	
ttt cag Phe Gln	atc cto	u Thr	gga Gly	gag Glu	gat Asp	tgg Trp 665	aat Asn	gcc Ala	gta Val	atg Met	tat Tyr 670	cat His	gly aaa	2136	
att gag Ile Glu														2184	
ttc atc Phe Ile 690	gtc cto	g aca u Thr	ctg Leu	ttt Phe 695	gga Gly	aac Asn	tac Tyr	acc Thr	ctg Leu 700	ctg Leu	aat Asn	gtt Val	ttt Phe	2232	
ctg gcc Leu Ala 705														2280	
gat gaa Asp Glu														2328	
aag gcc Lys Ala														2376	

tcc atc gct gcg cag gag aac tcg gcc aag gcg cgc tca gta tgg gag 2424 Ser Ile Ala Ala Gln Glu Asn Ser Ala Lys Ala Arg Ser Val Trp Glu 760 cag egg gee agt cag eta agg ete cag aat etg egt gee age tgt gag 2472 Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys Glu 775 780 gca ttg tac agt gag atg gac cct gag gag cgc ctg cgt tat gcc agc 2520 Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Tyr Ala Ser 790 acg cgc cat gtg agg cca gac atg aag aca cac atg gac cga ccc cta 2568 Thr Arg His Val Arg Pro Asp Met Lys Thr His Met Asp Arg Pro Leu 810 gtg gtg gag cct ggt cga gat ggc ttg cgg gga ccc gtt ggg agc aag 2616 Val Val Glu Pro Gly Arg Asp Gly Leu Arg Gly Pro Val Gly Ser Lys tca aag cct gaa ggc acg gag gcc aca gaa agc gcg gac cta cct cgc 2664 Ser Lys Pro Glu Gly Thr Glu Ala Thr Glu Ser Ala Asp Leu Pro Arg 840 cgg cac cac cgg cat cgt gat agg gac aag acc tca gcc aca gca cct 2712 Arg His Arg His Arg Asp Arg Asp Lys Thr Ser Ala Thr Ala Pro get gga gge gaa cag gac agg aca gaa age ace gag ace ggg cee egg 2760 Ala Gly Gly Glu Gln Asp Arg Thr Glu Ser Thr Glu Thr Gly Pro Arg 865 870 gag gaa cgt gcg cgc cct cgt cga agt cac agc aag gag act cca qqq 2808 Glu Glu Arg Ala Arg Pro Arg Arg Ser His Ser Lys Glu Thr Pro Gly 885 890 get gac acg caa gtg cgc tgt gag cgc agt agg cgt cac cac cgg cgc 2856 Ala Asp Thr Gln Val Arg Cys Glu Arg Ser Arg Arg His His Arg Arg 900 905 gge tee eeg gag gag gee aet gaa egg gag eet egg ege eac egt gee 2904 Gly Ser Pro Glu Glu Ala Thr Glu Arg Glu Pro Arg Arg His Arg Ala 920 cac cgg cat gca cag gac tca agc aag gag ggc acg gcg ccg gtg ctt 2952 His Arg His Ala Gln Asp Ser Ser Lys Glu Gly Thr Ala Pro Val Leu 935 gta ccc aag ggt gag cga cga gca aga cac cga ggc cca cgc acg ggt 3000 Val Pro Lys Gly Glu Arg Arg Ala Arg His Arg Gly Pro Arg Thr Gly 950 cca cgt gag gca gag aac aac gag gag ccc aca cgc agg cac cgt gca 3048 Pro Arg Glu Ala Glu Asn Asn Glu Glu Pro Thr Arg Arg His Arg Ala 965

		Lys '				aca ct Thr Le	eu G						ı Ālā		3096	
	Lys (					Val G					Glu Th			aat cac Asn His	3144	
Gln		Lys				tgt Cys 1015	Asp								3189	
Gly						atg Met 1030									3234	
Val	_					gat Asp 1045	_	_		_	_		_		3279	
Arg		Gly				tca Ser 1060									3324	
Pro			_			cct Pro 1075						_	_		3369	
Ser		Asn				gaa Glu 1090									3414	
Ala	gag Glu 1100		_	_		ctg Leu 1105	_	_						-	3459	
Pro	tac Tyr 1115	_		_		tgt Cys 1120		_				_		_	3504	
Arg	Phe	Cys	His	Tyr	Ile	gtg Val 1135	Thr	Met	Arg	Tyr	Leu	Glu			3549.	
Ile						ttg Leu 1150									3594	
Asp						tca Ser 1165									3639 .	
		Tyr				gga Gly 1180									3684	

(

	atg Met 1190					ttg Leu 1195									3729
	gac Asp 1205					ctg Leu 1210									3774
						tca Ser 1225									3819
						gtc Val 1240									3864
	atc Ile 1250					aaa Lys 1255									3909
	aac Asn 1265					gtc Val 1270									3954
	ttc Phe 1280					gcc Ala 1285									3999
						act Thr 1300									4044
	_			_		ttg Leu 1315	_			_	_	_	_		4089
						aag Lys 1330									4134
Val	Leu	${\tt Trp}$	Ala	Leu	Leu	acg Thr 1345	Leu					Thr			4179
						aaa Lys 1360									4224
						ggc Gly 1375									4269
tac Tyr	gtg Val 1385	gtc Val	tac Tyr	ttt Phe	gtg Val	gtc Val 1390	ttc Phe	cct Pro	ttt Phe	ttc Phe	ttt Phe 1395	gtc Val	aac Asn	atc Ile	4314
ttt	gtg	gcc	ttg	atc	att	atc	acc	ttc	cag	gaa	cag	gga	gat	aag	4359

Phe	Val 1400	Ala	Leu	Ile	Ile	Ile 1405	Thr	Phe	Gln	Glu	Gln 1410	Gly	Asp	Lys			
						tta Leu 1420									4404		
_		_		-	-	aag Lys 1435		_				_			4449		
		Gln				tat Tyr 1450	Lys								4494		
						atc Ile 1465									4539		
						ttc Phe 1480									4584		
						aac Asn 1495									4629		
						atc Ile 1510									4674		
						gtc Val 1525									4719	r v	
agt Ser	att Ile 1535	act Thr	gat Asp	att Ile	tta Leu	gta Val 1540	aca Thr	gag Glu	att Ile	gcg Ala	gaa Glu 1545	acg Thr	aac Asn	aac Asn	4764		
ttc Phe	atc Ile 1550	aac Asn	cta Leu	agc Ser	ttc Phe	ctt Leu 1555	cgc Arg	ctc Leu	ttc Phe	cgg Arg	gcg Ala 1560	gca Ala	cgg Arg	ctg Leu	4809		
						ggc Gly 1570									4854		
						aag Lys 1585									4899		
						atc Ile 1600									4944		
						gat Asp						aac Asn			4989		

	1610					1615					1620				
						ctg Leu 1630									5034
_	gcc Ala 1640					tgg Trp 1645	His								5079
_						cca Pro 1660									5124
-	_		_			tat , Tyr 1675		_						_	5169
		_	_	_		ctc Leu 1690		_	_	_	atc Ile 1695	_	_		5214
						gac Asp 1705									5259
						gtc Val 1720					gac Asp 1725				5304
						aat Asn 1735									5349
	tcc Ser 1745					ttg Leu 1750									5394
_			_	_	_	cgc Arg 1765	_		_		ata Ile 1770				5439
						acg Thr 1780									5484
						ctt Leu 1795									5529
						ctc Leu 1810									5574
	aaa Lys 1820			_	_	cag Gln 1825					ccc Pro 1830			~	5619

		Met				tcc Ser 1840	Leu							acc Thr	5664
						gct Ala 1855									5709
		Gln				act Thr 1870	Ser								5754
caa Gln	acc Thr 1880	cag Gln	gaa Glu	agt Ser	gga Gly	tca Ser 1885	agg Arg	agt Ser	cgc Arg	tgt Cys	cct Pro 1890	Gly 399	gga Gly	cgc Arg	5799
						ctt Leu 1900									5844
cgt Arg	gac Asp 1910	cat His	tct Ser	aaa Lys	gag Glu	atc Ile 1915	cct Pro	gtg Val	Gly ggg	cag Gln	tca Ser 1920	gga Gly	aca Thr	ctg Leu	5889
ctg Leu	gtg Val 1925	gat Asp	gtc Val	cag Gln	atg Met	cag Gln 1930	ạac Asn	atg Met	aca Thr	ctg Leu	aga Arg 1935	gga Gly	cca Pro	gat Asp	5934
Gly aaa	gat Asp 1940	ccc Pro	cag Gln	cct Pro	ggc Gly	ctg Leu 1945	gaa Glu	agc Ser	caa Gln	ggc Gly	aga Arg 1950	gct Ala	gcc Ala	tct Ser	5979
						gaa Glu 1960							gcc Ala	_	6024
ccc Pro	atg Met 1970	aag Lys	cgc Arg	tcc Ser	atc Ile	tcc Ser 1975	aca Thr	ctg Leu	gct Ala	cca Pro	cgc Arg 1980	cca Pro	gat Asp	gjå aaa	6069
Thr	Gln	Leu	Cys	Ser	Thr	gtt Val 1990	Leu	Asp	Arg	Pro	cct Pro 1995	Pro	agc Ser	cag Gln	6114
						cac His 2005									6159
_						aag Lys 2020									6204
	ggt Gly 2030	gca Ala	cca Pro	agc Ser	act Thr	gct Ala 2035	gct Ala	gca Ala	gga Gly	cct Pro	ggt Gly 2040	ctg Leu	ccc Pro	cat His	6249

											aaa Lys 2055				6294
							Gln				tcc Ser 2070				6339
											gcc Ala 2085				6384
											tcg Ser 2100				6429
				_				_		_	ggt Gly 2115		_		6474
											act Thr 2130				6519
				_			_			_	acc Thr 2145		_		6564
_				_		_			_		gtc Val 2160			_	6609
											ggc Gly 2175				6654
											aaa Lys 2190			ctg Leu	6699
											tct Ser 2205				6744
		_	_	_	_	_		_		_	cac His 2220		_		6789
											acc Thr 2235				6834
_	tcc Ser 2240										act Thr 2250				6879
cac	cct	ctc	cgc	cgt	gta	ccc	aat	ggc	tat	cac	tgc	act	ttg	gga	6924

His Pro Leu Arg Arg Val Pro Asn Gly Tyr His Cys Thr Leu Gly 2255 2260 2265	
ctc aac act ggc gtc ggg gca cga gca agc tac cac cac ccc gat Leu Asn Thr Gly Val Gly Ala Arg Ala Ser Tyr His His Pro Asp 2270 2275 2280	6969
cag gac cac tgg tgc tag cttcaccacg accacccatg taccagctcc Gln Asp His Trp Cys 2285	7017
atgggtgagg gttccagttg atgagtttta tcatcccact ctggactgtg gggtcacaac	7077
cetgggagga gggccetcac ateteggeet etgtggtgga ggeteetget teceteeete	7137
cctccctttt tacactggat agactaataa agccctttct tagagggg	7185
<210> 40 <211> 2288 <212> PRT <213> Mus musculus	
<400> 40	
Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Thr Gly 1 5 10 15	
Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro 20 25 . 30	
Gly Gln Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45	
Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50 55 60	
Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80	
Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 95	
Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110	
Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125	

arg	130	Asp	Asp	THE	GIU	135	Tyr	Pne	пе	GIY	11e 140	Pne	Cys	Pne	GIU
Ala 145	Gly	Ile	Lys	Ile	Ile 150	Ala	Leu	Gly	Phe	Val 155	Phe	His	Lys	Gly	Ser 160
Tyr	Leu	Arg	Asn	Gly 165	Trp	Asn	Val	Met	Asp 170	Phe	Val	Val	Val	Leu 175	Thr
Gly	Ile	Leu	Ala 180	Thr	Ala	Gly	Thr	Asp 185	Phe	Asp	Leu	Arg	Thr 190	Leu	Arg
Ala	Val	Arg 195	Val	Leu	Arg	Pro	Leu 200	Lys	Leu	Val	Ser	Gly 205	Ile	Pro	Ser
Leu	Gln 210	Val	Val	Leu	Lys	Ser 215	Ile	Met	Lys	Ala	Met 220	Val	Pro	Leu	Leu
Gln 225	Ile	Gly	Leu	Leu	Leu 230	Phe	Phe	Ala	Ile	Leu 235	Met	Phe	Gly	Ile	Ile 240
Gly	Leu	Glu	Phe	Tyr 245	Met	Gly	Lys	Phe	His 250	Lys	Ala	Cys	Phe	Pro 255	Asn
Ser	Thr	Asp	Thr 260	Glu	Pro	Val	Gly	Asp 265	Phe	Pro	Cys	Gly	Lys 270	Asp	Pro
Pro	Ala	Arg 275	Gln	Cys	Asp	Gly	Asp 280	Thr	Glu	Сув	Arg	Glu 285	Tyr	Trp	Pro
Gly	Pro 290	Asn	Phe	Gly	Ile	Thr 295	Asn	Phe	Asp	Asn	Ile 300	Leu	Phe	Ala	Ile
Leu 305	Thr	Val	Phe	Gln	Cys 310	Ile	Thr	Met	Glu	Gly 315	Trp	Thr	Asp	Ile	Leu 320
Tyr	Asn	Thr	Asn	Asp 325	Ala	Ala	Gly	Asn	Thr 330	Trp	Asn	Trp	Leu	Tyr 335	Phe
- 1	_	_				~3	_	_,	_,		_	_	_		_

Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn

350

340

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu

355 360 365

Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Ile Glu Arg Glu 375 370 Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 390 395 400 Ala Glu Glu Asp Lys Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu 405 410 415 Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu 425 Glu Gly Glu Asp Arg Phe Val Asp Leu Cys Ala Val Gly Ser Pro Phe Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe 450 455 Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys 470 475 Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Gln Arg Leu Thr Thr 500 505 Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe Glu 550 555 Val Val Trp Ala Ala Ile Lys Pro Gly Thr Ser Phe Gly Ile Ser Val 570 Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp

585

- Asn Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 595 600 605
- Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala 610 615 620
- Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 635 640
- Thr Pro Thr Thr Ile Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val 645 650 655
- Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly 660 665 670
- Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr 675 680 685
- Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe 690 695 700
- Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys 705 710 715 720
- Asp Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln 725 730 735
- Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750
- Ser Ile Ala Ala Gln Glu Asn Ser Ala Lys Ala Arg Ser Val Trp Glu 755 760 765
- Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys Glu 770 780
- Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Tyr Ala Ser 785 790 795 800
- Thr Arg His Val Arg Pro Asp Met Lys Thr His Met Asp Arg Pro Leu 805 810 815

- Val Val Glu Pro Gly Arg Asp Gly Leu Arg Gly Pro Val Gly Ser Lys 820 825 830
- Ser Lys Pro Glu Gly Thr Glu Ala Thr Glu Ser Ala Asp Leu Pro Arg 835 840 845
- Arg His His Arg His Arg Asp Arg Asp Lys Thr Ser Ala Thr Ala Pro 850 855
- Ala Gly Glu Glu Asp Arg Thr Glu Ser Thr Glu Thr Gly Pro Arg 865 870 875 886
- Glu Glu Arg Ala Arg Pro Arg Arg Ser His Ser Lys Glu Thr Pro Gly 885 890 895
- Ala Asp Thr Gln Val Arg Cys Glu Arg Ser Arg Arg His His Arg Arg 900 905 910
- Gly Ser Pro Glu Glu Ala Thr Glu Arg Glu Pro Arg Arg His Arg Ala 915 920 925
- His Arg His Ala Gln Asp Ser Ser Lys Glu Gly Thr Ala Pro Val Leu 930 935 940
- Val Pro Lys Gly Glu Arg Arg Ala Arg His Arg Gly Pro Arg Thr Gly 945 955 960
- Pro Arg Glu Ala Glu Asn Asn Glu Glu Pro Thr Arg Arg His Arg Ala . 965 970 975
- Arg His Lys Val Pro Pro Thr Leu Gln Pro Pro Glu Arg Glu Ala Ala 980 985 990
- Glu Lys Glu Ser Asn Ala Val Glu Gly Asp Lys Glu Thr Arg Asn His 995 1000 1005
- Gln Pro Lys Glu Pro His Cys Asp Leu Glu Ala Ile Ala Val Thr 1010 1015 1020
- Gly Val Gly Pro Leu His Met Leu Pro Ser Thr Cys Leu Gln Lys 1025 1030 1035

- Val Asp Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr 1040 1045 1050
- Arg Met Gly Ser Gln Pro Ser Asp Pro Ser Thr Thr Val His Val 1055 1060 1065
- Pro Val Thr Leu Thr Gly Pro Pro Gly Glu Thr Pro Val Val Pro 1070 1075 1080
- Ser Gly Asn Met Asn Leu Glu Gly Gln Ala Glu Gly Lys Lys Glu 1085 1090 1095
- Ala Glu Ala Asp Asp Val Leu Arg Arg Gly Pro Arg Pro Ile Val 1100 1105 1110
- Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Phe Arg 1115 1120 1125
- Arg Phe Cys His Tyr Ile Val Thr Met Arg Tyr Leu Glu Met Val 1130 1135 1140
- Ile Leu Val Val Ile Ala Leu Ser'Ser Ile Ala Leu Ala Ala Glu 1145 1150 1155
- Asp Pro Val Arg Thr Asp Ser Phe Arg Asn Asn Ala Leu Glu Tyr 1160 1165 1170
- Met Asp Tyr Ile Phe Thr Gly Val Phe Thr Cys Glu Met Val Ile 1175 1180 1185
- Lys Met Ile Asp Leu Gly Leu Leu His Pro Gly Ala Tyr Phe 1190 1195 1200
- Arg Asp Leu Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala 1205 1210 1215
- Leu Val Ala Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn 1220 1225 1230
- Thr Ile Lys Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys 1235 1240 1245
- Thr Ile Lys Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val

1250	1255	1260

Val	Asn 1265	Ser	Leu	Lys	Asn	Val 1270		Asn	Ile	Leu	Ile 1275	Val	Tyr	Met
Leu	Phe 1280	Met	Phe	Ile	Phe	Ala 1285	Val	Ile	Ala	Val	Gln 1290	Leu	Phe	Lys
Gly	Lys 1295		Phe	Tyr	Cys	Thr 1300	Asp	Glu	Ser	Lys	Glu 1305	Leu	Glu	Arg
Asp	Cys 1310	Arg	Gly	Gln	Tyr	Leu 1315	Asp	Tyr	Glu	Lys	Glu 1320	Glu	Val	Glu
Ala	Gln 1325	Pro	Arg	Gln	Trp	Lys 1330	Lys	Tyr	Asp	Phe	His 1335	Tyr	Asp	Asn
Val	Leu 1340	Trp	Ala	Leu	Leu	Thr 1345	Leu	Phe	Thr	Val	Ser 1350	Thr	Gly	Glu
Gly	Trp 1355	Pro	Met	Val	Leu	Lys 1360	His	Ser	Val	Asp	Ala 1365	Thr	Tyr	Glu
Glu	Gln 1370	Gly	Pro	Ser	Pro	Gly 1375	Phe	Arg	Met	Glu	Leu 1380	Ser	Ile	Leu
	Val 1385					1390					1395			
Phe	Val 1400	Ala	Leu	Ile	Ile	Ile 1405	Thr	Phe	Gln	Glu	Gln 1410	Gly	Asp	Lys
Val	Met 1415	Ser	Glu	Cys		Leu 1420		Lys	Asn	Glu	Arg 1425	Ala	Cys	Ile
	Phe 1430					1435				_	1440			
Asn	Lys 1445	Gln	Ser	Phe	Gln	Tyr 1450	Lys	Thr	Trp	Thr	Phe 1455	Val	Val	Ser
Pro	Pro 1460	Phe	Glu	Tyr	Phe	Ile 1465	Met	Ala	Met	Ile	Ala 1470	Leu	Asn	Thr

.

- Val Val Leu Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu 1475 1480 1485
- Leu Met Leu Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser 1490 1495 1500
- Met Glu Cys Ile Leu Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr 1505 1510 1515
- Phe Arg Asp Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly 1520 1530
- Ser Ile Thr Asp Ile Leu Val Thr Glu Ile Ala Glu Thr Asn Asn 1535 1540 1545
- Phe Ile Asn Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu 1550 1560
- Ile Lys Leu Leu Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp 1565 1570 1575
- Thr Phe Val Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu 1580 1585 1590
- Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val 1595 1600 1605
- Phe Gly Asn Ser Ala Leu Asp Asp Asp Thr Ser Ile Asn Arg His 1610 1615 1620
- Asn Asn Phe Arg Thr Phe Leu Gln Ala Ile Met Leu Leu Phe Arg 1625 1630 1635
- Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys Leu 1640 1645 1650
- Asp Asn Arg Ala Cys Asp Pro His Ala Asn Ala Ser Glu Cys Gly 1655 1660 1665
- Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys 1670 1675 1680

- Ser Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn 1685 1690 1695
- Phe Glu Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His 1700 1705 1710
- Leu Asp Glu Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala 1715 1720 1725
- Cys Gly Arg Ile Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His 1730 1740
- Met Ser Pro Pro Leu Gly Leu Gly Lys Lys Cys Pro Ala Arg Val 1745 1750 1755
- Ala Tyr Lys Arg Leu Val Arg Met Asn Met Pro Ile Ser Asn Glu 1760 1765 1770
- Asp Met Thr Val His Phe Thr Ser Thr Leu Met Ala Leu Ile Arg 1775 1780 1785
- Thr Ala Leu Glu Ile Lys Leu Ala Pro Ala Asp Glu Met Thr Val 1790 1795 1800
- Gly Lys Val Tyr Ala Ala Leu Met Ile Phe Asp Phe Tyr Lys Gln 1805 1810 1815
- Asn Lys Thr Thr Arg Asp Gln Thr His Gln Ala Pro Gly Gly Leu 1820 1825 1830
- Ser Gln Met Gly Pro Val Ser Leu Phe His Pro Leu Lys Ala Thr 1835 1840 1845
- Leu Glu Gln Thr Gln Pro Ala Val Leu Arg Gly Ala Arg Val Phe 1850 1855 1860
- Leu Arg Gln Lys Ser Ala Thr Ser Leu Ser Asn Gly Gly Ala Ile 1865 1870 1875
- Gln Thr Gln Glu Ser Gly Ser Arg Ser Arg Cys Pro Gly Gly Arg 1880 1885 1890

- Arg Gly Thr Gln Asp Ala Leu Tyr Glu Gly Arg Ala Pro Leu Glu 1895 1900 1905
- Arg Asp His Ser Lys Glu Ile Pro Val Gly Gln Ser Gly Thr Leu 1910 1915 1920
- Leu Val Asp Val Gln Met Gln Asn Met Thr Leu Arg Gly Pro Asp 1925 1930 1935
- Gly Asp Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg Ala Ala Ser 1940 1945 1950
- Met Leu Arg Leu Ala Ala Glu Thr Gln Pro Ala Pro Asn Ala Ser 1955 1960 1965
- Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Pro Arg Pro Asp Gly 1970 1975 1980
- Thr Gln Leu Cys Ser Thr Val Leu Asp Arg Pro Pro Pro Ser Gln 1985 1990 1995
- Ala Ser His His His His Arg Cys His Arg Arg Arg Asp Lys 2000 2010
- Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu Ser Val Asp Pro 2015 2020 2025
- Glu Gly Ala Pro Ser Thr Ala Ala Gly Pro Gly Leu Pro His 2030 2035 2040
- Gly Glu Gly Ser Thr Ala Cys Arg Arg Asp Arg Lys Gln Glu Arg 2045 2050 2055
- Gly Arg Ser Gln Glu Arg Arg Gln Pro Ser Ser Ser Ser Glu 2060 2065 2070
- Lys Gln Arg Phe Tyr Ser Cys Asp Arg Leu Gly Ala Gly Ser Pro 2075 2080 2085
- Gln Leu Met Pro Ser Leu Ser Ser His Pro Thr Ser Pro Ala Ala 2090 2095 2100
- Ala Leu Glu Pro Ala Pro His Pro Gln Gly Ser Gly Ser Val Asn

2110 2105 2115

Gly	Ser 2120	Pro	Leu	Met	Ser	Thr 2125		Gly	Ala	Ile	Thr 2130	Pro	Gly	Arg
Gly	Gly 2135	Arg	Arg	Gln	Leu	Pro 2140	Gln	Thr	Pro	Leu	Thr 2145	Pro	Arg	Pro
Ser	Ile 2150		Tyr	Lys	Thr	Ala 2155		Ser	Ser	Pro	Val 2160	His	Phe	Ala
Glu	Gly 2165	Gln	Ser	Gly	Leu	Pro 2170	Ala	Phe	Ser	Pro	Gly 2175	Arg	Leu	Ser
Arg	Gly 2180	Leu	Ser	Glu	His	Asn 2185	Ala	Leu	Leu	Gln	Lys 2190	Glu	Pro	Leu
Ser	Gln 2195		Leu	Ala	Pro	Gly 2200		Arg	Ile	Gly	Ser 2205	Asp	Pro	Tyr
Leu	Gly 2210	Gln	Arg	Leu	Asp	Ser 2215	Glu	Ala	Ser	Ala	His 2220	Thr	Leu	Pro
Glu	Asp 2225	Thr	Leu	Thr	Phe	Glu 2230	Glu	Ala	Val	Ala	Thr 2235	Asn	Ser	Gly

His Pro Leu Arg Arg Val Pro Asn Gly Tyr His Cys Thr Leu Gly 2255 2260

Arg Ser Ser Arg Thr Ser Tyr Val Ser Ser Leu Thr Ser Gln Ser 2240 2245 2250

Leu Asn Thr Gly Val Gly Ala Arg Ala Ser Tyr His His Pro Asp 2270 2275

Gln Asp His Trp Cys 2285

<210> 41 <211> 7713 <212> DNA

<213> Oryctolagus cuniculus

<220>

<221> CDS <222> (121)..(7140)

<400> 41 ggccgggccg ggcgg	ggtcc ggtgcggtc	c ggcagctccg	cggcggcttc gtc	tcgatcg 60
cetggegege cetee	ctgcc ggagccgcc	g ggccggggat	gcgcgcggcg ccc	ggtggcc 120
atg gtc cgc ttc of Met Val Arg Phe of				
ggc gcg gag cgg g Gly Ala Glu Arg 20				<del>-</del>
ggc ccc ggg ggg Gly Pro Gly Gly 35	ctg ccg ccg ggc Leu Pro Pro Gly 40	cag cgg gtc Gln Arg Val	ctc tac aaa ca Leu Tyr Lys Gl 45	g tcg 264 n Ser
atc gcg cag cgc g Ile Ala Gln Arg 7				
aag cag aac tgc Lys Gln Asn Cys 65				
gac aac gtc gtc Asp Asn Val Val	_			_
ttc gag tac atg Phe Glu Tyr Met 100			•	
gcc ctg gag cag o Ala Leu Glu Gln 1 115				
cgc ctg gat gac a Arg Leu Asp Asp '				
gcg ggg atc aag Ala Gly Ile Lys 145				
tat ctg cgg aac o Tyr Leu Arg Asn o				u Thr
ggg att ctt gcc a Gly Ile Leu Ala 1 180				

gct Ala																744
Leu				ctc Leu												792
_			_	ctg Leu				_			_		_			840
ggc Gly																888
				gat Asp												936
				tgt Cys												984
Gly				ggt Gly					_			_		_		1032
				cag Gln												1080
				gat Asp 325												1128
				atc Ile						_			_		_	1176
		Leu	Ser	gga Gly	Glu	Phe	Ala	Lys	Glu	Arg	Glu	Arg	Val			1224
Arg	_			ctg Leu	_				_	_	_					1272
ctc Leu 385																1320
gct Ala			_			_			_			_	_	_		1368
ttg	aag	aga	gca	gct	gcc	aag	aag	agc	aga	agc	gac	ctg	atc	cag	gcc	1416

Leu Lys	Arg Al		Ala	Lys	Lys	Ser 425	Arg	Ser	Asp	Leu	Ile 430	Gln	Ala		
gag gag Glu Glu								_	_				_	1464	
ttc gcc Phe Ala 450		_	_		_		_	_		_	_	_		1512	
ttc cgg Phe Arg 465			_	_		_					_	_		1560	
aag gca Lys Ala														1608	
aca ctg Thr Leu		l Ala												1656	
acg gca Thr Ala	_		_			_		-	~ ~					1704	
gag atg Glu Met 530		_	_			_				_				1752	
tcc tcc Ser Ser 545														1800	*, *
gaa gtg Glu Val														1848	
gtt ctg Val Leu		a Leu	_		_	-			_	_				1896	
tgg aac Trp Asn														1944	
tcc atc Ser Ile 610					_			_			_	_		1992	
gct ctg Ala Leu 625														2040	
gag acc Glu Thr														2088	

•

645 650 655

							gac Asp 665					:	2136
			_		 	_	agc Ser	_	 _			:	2184
							GJA aaa					:	2232
	_	_		-	 _		ctt Leu	_	_	_	 _	:	2280
							gca Ala					:	2328
							gtc Val 745					:	2376
							aac Asn					:	2424
							cgg Arg					:	2472
							gac Asp					:	2520
							gac Asp					:	2568
							gac Asp 825					:	2616
							gag Glu					:	2664
							gac Asp					:	2712
							aga Arg					:	2760

ggc (	31y	gag Glu	ctg Leu	999 Gly 885	ccc Pro	cgc Arg	gag Glu	gag Glu	cgg Arg 890	Gly 999	cgg Arg	cca Pro	cgc Arg	cgc Arg 895	agc Ser	2808
cgc a																2856
cgt ( Arg (																2904
ccg g Pro (																2952
cac q His ( 945																3000
gag ( Glu <i>l</i>																3048
gct o																3096
gcg ( Ala I	ccg Pro	ccc Pro 995	acc Thr	cag Gln	gag Glu	acc Thr	gcg Ala 1000	Glı	g aaq ı Ly:	g gad s Ası	c aaq o Lys	g gaq s Gli 100	u Al	et go la Al	cc gag la Glu	3144
aag g Lys (		Gly					ı <sub>.</sub> A]				sp Ly		gag g Glu <i>l</i>			3189
aac ( Asn H		Gln		_		~	Pr	_	_		eu G.		gcc a Ala 1	_	, _	3234
atg d Met I	_	Gly			_		Hi		_	_	o Se	_	acc t Thr (	_	_	3279
cag a Gln I		Val					G1				sp As					3324
gtc a Val 1		Arg					Pr				ır Se					3369
cac a		Pro					G1				y G		acc a Thr T			3414

gtg Val	ccc Pro 1100	agt Ser	gga Gly	aac Asn	gtg Val	gac Asp 1105	Leu	gaa Glu	agt Ser	caa Gln	gca Ala 1110	gag Glu	G1 y 999	aag Lys	3459
aag Lys	gag Glu 1115	gtg Val	gaa Glu	acc Thr	agc Ser	gat Asp 1120	gtg Val	atg Met	agg Arg	agc Ser	ggc Gly 1125	cct Pro	cgg Arg	ccc Pro	3504
	gtc Val 1130					atg Met 1135	Phe								3549
						tac Tyr 1150									3594
	gtc Val 1160	atc Ile	ctc Leu	gtg Val	gtc Val	atc Ile 1165	gcc Ala	ctg Leu	agc Ser	agc Ser	atc Ile 1170	gcc Ala	ctg Leu	gct Ala	3639
gcc Ala	gag Glu 1175	gac Asp	ccc Pro	gtg Val	aga Arg	aca Thr 1180	gac Asp	tct Ser	ccc Pro	agg Arg	aat Asn 1185	aat Asn	gct Ala	ctg Leu	3684
						ttc Phe 1195									3729
						ttg Leu 1210									3774
						aac Asn 1225									3819
Gly	gcc Ala 1235	ctg Leu	gtg Val	gca Ala	ttt Phe	gct Ala 1240	ttc Phe	tca Ser	gga Gly	tcc Ser	aaa Lys 1245	Gly aaa	aaa Lys	gac Asp	3864
Ile	Ser	Thr	Ile	Lys	Ser	ctg Leu 1255	Arg	Val	Leu	Arg	Val	Leu	Arg	cca Pro	3909
ctc Leu	aag Lys 1265	acc Thr	atc Ile	aag Lys	cgg Arg	ctg Leu 1270	ccc Pro	aag Lys	ctc Leu	aag Lys	gcc Ala 1275	gtg Val	ttt Phe	gac Asp	3954
						aaa Lys 1285									3999
Tyr	atg Met 1295	ctc Leu	ttc Phe	atg Met	ttc Phe	atc Ile 1300	ttt Phe	gcc Ala	gtc Val	att Ile	gcc Ala 1305	gtg Val	cag Gln	ctc Leu	4044
ttc	aaa	999	aag	ttc	ttc	tac	tgc	acg	gac	gaa	tcc	aag	gag	ctg	4089

]	Phe	Lys 1310	-	Lys	Phe	Phe	Tyr 1315	Cys	Thr	Asp	Glu	Ser 1320	-	Glu	Leu			
			Asp				cag Gln 1330	Tyr								4134		
							cag Gln 1345	Trp								4179		
							ctg Leu 1360									4224		
							gtg Val 1375									4269		
							agc Ser 1390	Pro								4314		
							ttt Phe 1405									4359		
							atc Ile 1420									4404	• · · · · · · · · · · · · · · · · · · ·	
							tgc Cys 1435									4449	s	
							agc Ser 1450								atg Met	4494		
	Pro						ttc Phe 1465	Gln								4539		
							tac Tyr 1480									4584		
					_	_	atg Met 1495	_			-		_			4629		
				~	_		tgc Cys 1510	_					_		_	4674		
			-		-		ctg Leu	_			_				_	4719		

	1520				1525			1530			
		_	_	_		_	_	ttt Phe 1545	_	_	 4764
	gga Gly 1550							att Ile 1560			4809
								gcc Ala 1575			4854
	aag Lys 1580							atc Ile 1590			4899
								gtg Val 1605			4944
	gcc Ala 1610							ggc Gly 1620			4989
								atc Ile 1635			5034 .
	aac Asn 1640							ttg Leu 1650			5079
_	gcc Ala 1655			_			_	ctg Leu 1665	_	_	5124
_	agc Ser 1670	 _	_	_			_	agc Ser 1680		_	 5169
_								atc Ile 1695			5214
	ttt Phe 1700							atc Ile 1710			5259
	gag Glu 1715				gac Asp 1720			999 Gly 1725	cct Pro		5304
	gat Asp 1730							gac Asp 1740			5349

	999 Gly 1745				agt Ser 1750									5394
_	tcc Ser 1760		_		ctg Leu 1765		_	_	_		_	_	_	5439
	tac Tyr 1775				cgc Arg 1780							agt Ser		5484
_	atg Met 1790				aca Thr 1795									5529
	gcg Ala 1805				ctg Leu 1810									5574
_	tgc Cys 1820				agg Arg 1825									5619
					ctg Leu 1840									5664
	gac Asp 1850	 _	_		ggc Gly 1855		_		_	_	_	_		5709
_	gac Asp 1865				aac Asn 1870			_	_	_	_	act Thr	_	5754
	gcc Ala 1880				tcc Ser 1885							ctg Leu		5799
	ccg Pro 1895				ctg Leu 1900									5844
	-	 _		_	cgg Arg 1915	_	_			_		_	~	5889
	ggc Gly 1925				acc Thr 1930							gaa Glu		5934
_	tct Ser 1940	 		-	agg Arg 1945	Thr	_	_			_		_	5979

-

						ggc Gly 1960									6024	•	
_						gta Val 1975									6069		
						gag Glu 1990									6114		
ggc Gly	cga Arg 2000	gcc Ala	gcc Ala	tcc Ser	atg Met	ccc Pro 2005	cgc Arg	ctg Leu	gcc Ala	gca Ala	gaa Glu 2010	aca Thr	cag Gln	ccc Pro	6159		
						atg Met 2020									6204		
						cgg Arg 2035									6249		
						ccg Pro 2050									6294		
						cag Gln 2065									6339	٨	
						999 Gly 2080									6384	<b>4</b> ,	
						ggc Gly 2095						cgg Arg			6429		
						ggc Gly 2110									6474		
						aag Lys 2125									6519		
						cca Pro 2140									6564		
						gcc Ala 2155									6609		
cag	999	agt	ggt	tcg	gtt	cac	999	agc	ccc	ctg	ctg	tcc	aca	tct	6654		

.

Gln	Gly 2165	`Ser	Gly	Ser	Val	His 2170	Gly	Ser	Pro	Leu	Leu 2175	Ser	Thr	Ser		
	gct Ala 2180														6699	
	ttg Leu 2195														6744	
	cct Pro 2210														6789	
	ccc Pro 2225	Gly													6834	
	cag Gln 2240														6879	
	ggc Gly 2255														6924	
	gcc Ala 2270														6969	
	gcc Ala 2285														7014	.*
	ctg Leu 2300						Pro								7059	
tac Tyr	cac His 2315	Cys	act Thr	ctg Leu	glà aaa	ctg Leu 2320	ggc Gly	Gly ggg	ggc Gly	ggc Gly	cgg Arg 2325	gcg Ala	cgg Arg	cgc Arg	7104	
	tgc Cys 2330	His									tag (	ccgct	gege	eg	7150	
														accgagg	7210	
														agcagac	7270	
gga				•			•				•			gcagggg	7330 7390	

gagcacctgt	gacccgtgtt	ggtgggagga	gttgcactgc	ctggaggcat	aggggagctt	7510
aggggtctcg	agggagccgg	gggaggagct	gaggttgctc	ctggtccctg	gtcgcggggc	7570
ggcaggtgga	cgctggacac	actcagatct	ctcagtccct	ccccgacccc	atcagcccca	7630
acagggacca	aacctggagg	acgtgccggc	tgctggtcta	aacagaacag	gaagaaaatg	7690
aggaaaccgg	aaaatcagag	ccg				7713

<210> 42

<211> 2339

<212> PRT

<213> Oryctolagus cuniculus

<400> 42

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Ala Gly 1 5 10 15

Gly Ala Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro
20 25 30

Gly Pro Gly Gly Leu Pro Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50 55 60

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140

- Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr
  165 170 175
- Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 185 190
- Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205
- Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 215 220
- Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile 225 230 235 240
- Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 255
- Ser Thr Asp Pro Asp Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala 260 265 270
- Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Ala 275 280 285
- Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 290 295 300
- Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 305 310 315 320
- Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 325 330 335
- Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 350
- Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365
- Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 375 380

- Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 390 395 400
- Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Ala Val405 410 415
- Leu Lys Arg Ala Ala Ala Lys Lys Ser Arg Ser Asp Leu Ile Gln Ala 420 425 430
- Glu Glu Gly Glu Gly Arg Leu Thr Gly Leu Cys Ala Pro Gly Ser Pro 435 440 445
- Phe Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Tyr 450 455 460
- Phe Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val 465 470 475 480
- Lys Ala Gln Ser Phe Tyr Trp Thr Val Leu Cys Val Val Ala Leu Asn 485 490 495
- Thr Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Gln Arg Leu Thr 500 505 510
- Thr Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr 515 520 525
- Glu Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg 530 540
- Ser Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe 545 550 555
- Glu Val Val Trp Ala Ala Val Lys Pro Gly Thr Ser Phe Gly Ile Ser 565 570 575
- Val Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr 580 585 590
- Trp Asn Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys 595 600 605
- Ser Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe

10	615 62	(

Ala 625	Leu	Leu	Gly	Met	Gln 630	Leu	Phe	Gly	Gly	Gln 635	Phe	Asn	Phe	Lys	Asp 640
Glu	Tḥr	Pro	Thr	Thr 645	Asn	Phe	Asp	Thr	Phe 650	Pro	Ala	Ala	Ile	Leu 655	Thr
Val	Phe	Gln	Ile 660	Leu	Thr	Gly	Glu	Asp 665	Trp	Asn	Ala	Val	Met 670	Tyr	His
Gly	Ile	Glu 675	Ser	Gln	Gly	Gly	Val 680	Ser	Arg	Gly	Met	Phe 685	Ser	Ser	Phe
Tyr	Phe 690	Ile	Val	Leu	Thr	Leu 695	Phe	Gly	Asn	Tyr	Thr 700	Leu	Leu	Asn	Val
Phe 705	Leu	Ala	Ile	Ala	Val 710	Asp	Asn	Leu	Ala	Asn 715	Ala	Gln	Glu	Leu	Thr 720
Lys	Asp	Glu	Glu	Glu 725	Met	Glu	Glu	Ala	Ala 730	Asn	Gln	Lys	Leu	Ala 735	Leu
Gln	Lys	Ala	Lys 740	Glu	Val	Ala	Glu	Val 745	Ser	Pro	Met	Ser	Ala 750	Ala	Asn
Ile	Ser	Ile 755	Ala	Ala	Arg	Gln	Gln 760	Asn	Ser	Ala	Lys	Ala 765	Arg	Ser	Val
Trp	Glu 770	Gln	Arg	Ala	Ser	Gln 775	Leu	Arg	Leu	Gln	Asn 780	Leu	Arg	Ala	Ser
Cys 785	Glu	Ala	Leu	Tyr	Ser 790	Glu	Met	Asp	Pro	Glu 795	Glu	Arg	Leu	Arg	Tyr 800
Ala	Thr	Ala	Arg	His 805	Leu	Arg	Pro	Asp	Val 810	Lys	Thr	His	Leu	Asp 815	Arg
Pro	Leu	Val	Val 820	Glu	Pro	Gly	Arg	Asp 825	Ala	Pro	Arg	Gly	Pro 830	Pro	Gly
Gly	Lys	Ser	Arg	Pro	Asp	Gly	Ser 840	Gl.u	Ala	Pro	Glu	Gly	Ala	Asp	Pro

- Pro Arg Arg His His Arg His Arg Asp Lys Asp Lys Ala Pro Ala Thr 850 855 860
- Val Pro Ser Ala Gly Glu Gln Asp Arg Ala Glu Ala Leu Arg Ala Glu 865 870 875 880
- Gly Gly Glu Leu Gly Pro Arg Glu Glu Arg Gly Arg Pro Arg Arg Ser 885 890 895
- Arg Ser Lys Glu Ala Pro Gly Ala Pro Glu Val Arg Ser Asp Arg Gly 900 905 910
- Arg Gly Pro Cys Pro Glu Gly Gly Arg Arg His His Arg Arg Gly Ser 915 920 925
- Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala His Arg 930 935 940
- His Gly Pro Asp Pro Gly Lys Glu Gly Pro Ala Ser Gly Thr Arg Gly 945 950 955 960
- Glu Arg Arg Ala Arg His Arg Thr Gly Pro Arg Ala Cys Pro Arg Glu 965 970 975
- Ala Glu Ser Ser Glu Glu Pro Ala Arg Arg His Arg Ala Arg His Lys 980 985 990
- Ala Pro Pro Thr Gln Glu Thr Ala Glu Lys Asp Lys Glu Ala Ala Glu 995 1000 1005
- Lys Gly Glu Ala Thr Glu Ala Glu Lys Asp Lys Glu Ala Arg 1010 1015 1020
- Asn His Gln Pro Lys Glu Leu Pro Cys Asp Leu Glu Ala Ile Gly 1025 1030 1035
- Met Leu Gly Val Gly Ala Val His Thr Leu Pro Ser Thr Cys Leu 1040 1045 1050
- Gln Lys Val Glu Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn 1055 1060 1065

- Val Thr Arg Met Gly Ser Gln Pro Pro Asp Thr Ser Thr Thr Val 1070 1075 1080
- His Ile Pro Val Thr Leu Thr Gly Pro Pro Gly Glu Thr Thr Val 1085 1090 1095
- Val Pro Ser Gly Asn Val Asp Leu Glu Ser Gln Ala Glu Gly Lys 1100 1110
- Lys Glu Val Glu Thr Ser Asp Val Met Arg Ser Gly Pro Arg Pro 1115 1120 1125
- Ile Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr Asn Leu 1130 1135 1140
- Leu Arg Arg Cys Cys His Tyr Ile Val Thr Met Arg Tyr Phe Glu 1145 1150 1155
- Met Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile Ala Leu Ala 1160 1165 1170
- Ala Glu Asp Pro Val Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu 1175 1180 1185
- Lys Tyr Met Asp Tyr Ile Phe Thr Gly Val Phe Thr Phe Glu Met 1190 1195 1200
- Val Ile Lys Met Ile Asp Leu Gly Leu Leu His Pro Gly Ala 1205 1210 1215
- Tyr Phe Arg Asp Leu Trp Asn Ile Leu Asp Phe Ile Val Val Ser 1220 1225 1230
- Gly Ala Leu Val Ala Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp 1235 1240 1245
- Ile Ser Thr Ile Lys Ser Leu Arg Val Leu Arg Val Leu Arg Pro 1250 1255 1260
- Leu Lys Thr Ile Lys Arg Leu Pro Lys Leu Lys Ala Val Phe Asp 1265 1270 1275

- Cys Val Val Asn Ser Leu Lys Asn Val Leu Asn Ile Leu Ile Val 1280 1285 1290
- Tyr Met Leu Phe Met Phe Ile Phe Ala Val Ile Ala Val Gln Leu 1295 1300 1305
- Phe Lys Gly Lys Phe Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu 1310 1315 1320
- Glu Arg Asp Cys Arg Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu 1325 1330 1335
- Val Glu Ala Gln Pro Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr 1340 1345 1350
- Asp Asn Val Leu Trp Ala Leu Leu Thr Leu Phe Thr Val Ser Thr 1355 1360 1365
- Gly Glu Gly Trp Pro Met Val Leu Lys His Ser Val Asp Ala Thr 1370 1375 1380
- Tyr Glu Glu Gln Gly Pro Ser Pro Gly Tyr Arg Met Glu Leu Ser 1385 1390 1395
- Ile Phe Tyr Val Val Tyr Phe Val Val Phe Pro Phe Phe Val 1400 1410

فد

- Asn Ile Phe Val Ala Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly 1415 1420 1425
- Asp Lys Val Met Ser Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala 1430 1440
- Cys Ile Asp Phe Ala Ile Ser Ala Arg Pro Leu Thr Arg Tyr Met 1445 1450 1455
- Pro Gln Asn Lys Gln Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val 1460 1465 1470
- Val Ser Pro Pro Phe Glu Tyr Phe Ile Met Ala Met Ile Ala Leu 1475 1480 1485
- Asn Thr Val Val Leu Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu

1490	1495	1500

Tyr	Glu 1505	Leu	Met	Leu	Lys	Cys 1510		Asn	Ile	Val	Phe 1515	Thr	Ser	Met
Phe	Ser 1520	Met	Glu	Cys	Val	Leu 1525	_	Ile	Ile	Ala	Phe 1530	Gly	Val	Leu
Asn	Туг 1535	Phe	Arg	Asp	Ala	Trp 1540	Asn	Val	Phe	Asp	Phe 1545	Val	Thr	Val
Leu	Gly 1550	Ser	Ile	Thr	Asp	Ile 1555		Val	Thr	Glu	Ile 1560		Asn	Asn
Phe	Ile 1565	Asn	Leu	Ser	Phe	Leu 1570	Arg	Leu	Phe	Arg	Ala 1575	Ala	Arg	Leu
Ile	Lys 1580	Leu	Leu	Arg	Gln	Gly 1585	Tyr	Thr	Ile	Arg	Ile 1590	Leu	Leu	Trp
Thr	Phe 1595	Val	Gln	Ser	Phe	Lys 1600	Ala	Leu	Pro	Tyr	Val 1605	Cys	Leu	Leu
Ile	Ala 1610	Met	Leu	Phe	Phe	Ile 1615	Tyr	Ala	Ile	Ile	Gly 1620	Met	Gln	Val
Phe	Gly 1625	Asn	Ile	Ala	Leu	Asp 1630	Asp	Asp	Thr	Ser	Ile 1635	Asn	Arg	His
Asn	Asn 1640	Phe	Arg	Thr	Phe	Leu 1645	Gln	Ala	Leu	Met	Leu 1650	Leu	Phe	Arg
Ser	Ala 1655	Thr	Gly	Glu	Ala	Trp 1660	His	Glu	Ile	Met	Leu 1665	Ser	Cys	Leu
Ser	Ser 1670	Arg	Ala	Cys	Asp	Glu 1675	His	Ser	Asn	Ala	Ser 1680	Glu	Cys	Gly
Ser	Asp 1685	Phe	Ala	Tyr	Phe	Tyr 1690	Phe	Val	Ser	Phe	Ile 1695	Phe	Leu	Cys
Ser	Phe 1700	Leu	Met	Leu	Asn	Leu 1705	Phe	Val	Ala	Val	Ile 1710	Met.	Asp	Asn

- Phe Glu Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His 1715 1720 1725
- Leu Asp Glu Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala 1730 1735 1740
- Cys Gly Arg Ile Ser Tyr Ser Asp Met Phe Glu Met Leu Lys His 1745 1750 1755
- Met Ser Pro Pro Leu Gly Leu Gly Lys Lys Cys Pro Ala Arg Val 1760 1765 1770
- Ala Tyr Lys Arg Leu Val Arg Met Asn Met Pro Ile Ser Ser Glu 1775 1780 1785
- Asp Met Thr Val His Phe Thr Ser Thr Leu Met Ala Leu Ile Arg 1790 1795 1800
- Thr Ala Leu Asp Ile Lys Leu Ala Pro Ala Gly Thr Lys Gln His 1805 1810 1815
- Gln Cys Asp Ala Glu Leu Arg Lys Glu Ile Ser Cys Val Trp Ala 1820 1825 1830 .
- Asn Leu Pro Gln Lys Thr Leu Asp Leu Leu Val Pro Pro His Lys 1835 1840 1845
- Pro Asp Glu Met Thr Val Gly Lys Val Tyr Ala Ala Leu Met Ile 1850 1860
- Phe Asp Phe Tyr Lys Gln Asn Lys Thr Ser Arg Asp Gln Thr Gln 1865 1870 1875
- Gln Ala Pro Gly Gly Leu Ser Gln Leu Gly Pro Val Ser Leu Phe 1880 1885 1890
- His Pro Leu Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Leu Arg 1895 1900 1905
- Gly Ala Arg Ala Phe Leu Arg Gln Lys Ser Ser Ala Ser Leu Ser 1910 1915 1920

- Asn Gly Gly Ala Val Gln Thr Gln Glu Ser Gly Ile Lys Glu Ser 1925 1930 1935
- Val Ser Trp Gly Thr Gln Arg Thr Gln Asp Val Leu Cys Glu Ala 1940 1945 1950
- Arg Ala Pro Leu Glu Arg Gly His Ser Ala Glu Ile Pro Val Gly 1955 1960 1965
- Gln Pro Gly Thr Leu Ala Val Asp Val Gln Met Gln Asn Met Thr 1970 1975 1980
- Leu Ser Gly Pro Asp Ala Glu Pro Gln Pro Gly Leu Glu Ser Gln 1985 1990 1995
- Gly Arg Ala Ala Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro 2000 2005 2010
- Ala Pro Asp Ala Ser Pro Met Lys Arg Ser Ile Ser Thr Leu Ala 2015 2020 2025
- Pro Arg Pro His Thr Ala Arg Leu Gly Ser Thr Ala Leu Asp Arg 2030 2035 2040
- Pro Ala Pro Ser Gln Ala Pro His His His His Arg Cys His 2045 2050 2055
- Arg Arg Asp Arg Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser 2060 2065 2070
- Leu Ser Ala Asp Thr Asp Gly Ala Pro Asp Ser Thr Val Gly Pro 2075 2080 2085
- Gly Leu Pro Thr Gly Glu Gly Pro Pro Gly Cys Arg Arg Glu Arg 2090 2095 2100
- Glu Arg Arg Gln Glu Arg Gly Arg Ser Gln Glu Arg Arg Gln Pro 2105 2110 2115
- Ser Ser Ser Ser Glu Lys His Arg Phe Tyr Ser Cys Asp Arg 2120 2125 2130

- Phe Gly Gly Arg Glu Pro Pro Gln Pro Lys Pro Ser Leu Ser Ser 2135 2140 2145
- His Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro His Pro 2150 2155 2160
- Gln Gly Ser Gly Ser Val His Gly Ser Pro Leu Leu Ser Thr Ser 2165 2170 2175
- Gly Ala Ser Thr Pro Gly Arg Gly Arg Gln Leu Pro Gln Thr 2180 2185 2190
- Pro Leu Thr Pro Arg Pro Ser Val Thr Tyr Lys Thr Ala Asn Ser 2195 2200 2205
- Ser Pro Val His Phe Ala Gly Ala Pro Ser Gly Leu Pro Ala Phe 2210 2215 2220
- Ser Pro Gly Arg Leu Ser Arg Gly Leu Ser Glu His Asn Ala Leu 2225 2230 2235
- Leu Gln Arg Asp Pro Leu Ser Arg Pro Leu Ala Pro Gly Ser Arg 2240 2245 2250
- Ile Gly Ser Asp Pro Tyr Leu Gly Gln Arg Leu Asp Ser Glu Ala 2255 2260 2265
- Pro Ala Arg Ala Leu Pro Glu Asp Ala Pro Ala Phe Glu Glu Thr 2270 2275 2280
- Ala Ala Ser Asn Ser Gly Arg Ser Ser Arg Thr Ser Tyr Val Ser 2285 2290 2295
- Ser Leu Thr Ser Gln Pro Pro Leu Arg Arg Val Pro Asn Gly 2300 2305 2310
- Tyr His Cys Thr Leu Gly Leu Gly Gly Gly Arg Ala Arg Arg 2315 2320 2325
- Gly Cys His His Pro Asp Arg Asp Arg Cys 2330 2335

7121 <211> <212> DNA <213> Bos taurus <220> <221> CDS <222> (126)..(7121) <400> 43 ctgggcgagg tctggtgcgg gtccggcggc tccgcggctg ctccgctaga gcgcagggcg 60 cacctegege etteegatee eegegggee geegggeegg gggatgeget eagegeeegg 120 gagec atg gtc ege ttc ggg gac gag etg ggc ege tat ggg ggc eec 170 Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro 218 Gly Gly Glu Arg Ala Arg Gly Gly Ala Gly Val Ala Gly Gly ccg ggc ccc ggg ggg ctg cag ccg ggc cag cgg gtc ctc tac aag cag 266 Pro Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln teg ate geg cag ege geg egg ace atg geg etg tae aac eec ate eeg 314 Ser Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro gte aag cag aac tgc ttc acc gtc aac cgc tcg ctc ttc gtc ttc agc 362 Val Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser gag gac aac qtc qtc cqc aaa tac qcc aaq cqc atc acc qag tqq cct 410 Glu Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro 85 90 ccg ttt gag tac atg att ctg gcc acc atc atc gcc aac tgt atc gtg 458 Pro Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val 100 105 ctg gct ctg gag cag cac ctc cca gac gga gac aag acg ccc atg tct 506 Leu Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser gag egg etg gac gac aeg gag eee tac tte ate gge ate tte tge tte 554 Glu Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe 135 gag gct ggg atc aag atc atc gcg ctg ggc ttt gtc ttg cac aag ggc 602 Glu Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Leu His Lys Gly 150

tcc tac ctg cgg aat ggc tgg aac gtc atg gac ttc gtg gtg gtc ctc

Ser Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Leu

170

165

650

_	Gly aaa		_		_			_		_	_	-		_	698
	gct Ala	 _	_	_			_	_	_						746
	ttg Leu														794
_	cag Gln 225		_					_			_		_		842
	ggc Gly				_					_	_	_			890
	agc Ser	_	_	_	_			_		_	-				938
	ccg Pro														986
	ggc Gly								-			_		_	1034
	ctg Leu 305														1082
	tac Tyr			_		_									1130
_	ctg Leu	_					_			_			_		1178
	ggc Gly														1226
	cgc Arg														1274
gag Glu	ctc								_	_			-	_	1322

.

ctg gcc Leu Ala 400			_	_		-			_			_	_		1370		
ttg aag Leu Lys		Ala i													1418		
gag gag Glu Glu	Gly														1466		
ttc gcc Phe Ala															1514		
ttc cgc Phe Arg 465	Arg I														1562		
aag gcg Lys Ala 480															1610		
acg ctg Thr Leu		Val i	_					_	_		_				1658	**	
acg gcg Thr Ala	Leu :			_			_		_						1706	::	
gag atg Glu Met															1754	<b>.</b>	
tcc tcc Ser Ser 545															1802		
gaa gtg Glu Val 560		Frp 7	Ala	Āla		Lys	Pro		Thr	Ser			Ile		1850		
gtg ctg Val Leu		Ala 1													1898		
tgg agc Trp Ser	Ser 1	_					~ ~		_				_	_	1946		
tcc atc Ser Ile															1994		
gct ctg	ctg (	ggc (	atg	cag	ctc	ttc	999	gga	cag	ttc	aac	ttc	cag	gat	2042		

Ala Leu Leu G 625	ly Met Gln Leu 630	Phe Gly Gly Gln	Phe Asn Phe Gln Asp 635	
		_	gcc gcc atc ctc act Ala Ala Ile Leu Thr 655	2090
			gcg gtg atg tac cat Ala Val Met Tyr His 670	2138
Gly Ile Glu S			atg ttc tcg tct gtg Met Phe Ser Ser Val 685	2186
_			acc ctg ctg aac gtc Thr Leu Leu Asn Val 700	2234
		Asn Leu Ala Asn	gct cag gag ctg acc Ala Gln Glu Leu Thr 715	2282
			cag aag ctt gct ctg Gln Lys Leu Ala Leu 735	2330
			atg tct gcc gcg aac Met Ser Ala Ala Asn 750	2378
Ile Ser Ile A			aag gcg cgc tcg gtg Lys Ala Arg Ser Val 765	2426 <u>.</u>
			aac ctg cgg gcc agc Asn Leu Arg Ala Ser 780	2474
			gag cgc ctg cgc ttc Glu Arg Leu Arg Phe 795	2522
			acg cac ctg gac cgg Thr His Leu Asp Arg 815	2570
			cgg ggg cct gcg ggg Arg Gly Pro Ala Gly 830	2618
Gly Lys Ala A			gag ggt gcc gac ccg Glu Gly Ala Asp Pro 845	2666
			agg gac aag gcc gca Arg Asp Lys Ala Ala	2714

•

850 855 860

	gcg ggg gag co Ma Gly Glu Pr					
	ccg ggg gcc cg Pro Gly Ala An 88	g Glu Glu		_		_
	gag gcc gca gg Slu Ala Ala Gl 900	_				_
	ggc ccg gac gg Gly Pro Asp Gl 915		_			_
Glu Glu A	gcc gcg gag cg Ala Ala Glu Ai 930			His Arg A		
	gag ccg agc ag Glu Pro Ser Ai					
	egg cac ege go Arg His Arg Gl 90	y Gly Ser				
	gag gag ccc to Glu Glu Pro Se 980					
_	cac gag gag go His Glu Glu Al 995			gag gcg a Glu Ala		gg gac 3146 rg Asp
Lys Glu P	ccc cga aac o Pro Arg Asn B 1010		o Arg Gl			_
	Ser Val Val	tt ggt gtg le Gly Vai	l Gly Pr		Ala Leu	
Ser Thr C	egt cta gag a Cys Leu Glu 1 1040		u Glu Gl			
Asn Gln A	egg aat gtg a Arg Asn Val 5 1055		Gly Se			
Ser Thr A	gct gtg cgt a Ala Val Arg : L070		l Thr L∈			

gag a Glu T													agc Ser		3416
gca g Ala G	Glu														3461
ggc c	Pro														3506
ccc a Pro T	rhr														3551
cag t Gln T	Гуr														3596
att g Ile A	Ala														3641
aac a Asn A	Asn														3686
acc t Thr P	Phe						atg Met 1195								3731
cac c His P	Pro														3776
att g Ile V	/al	_					gtg Val 1225								3821
aaa g Lys G	Gly												ctg Leu	_	3866
gtg c Val L	Leu														3911
gct g Ala V	/al	Phe 1265	Asp	Cys	Val	Val	1270	Ser	Leu	Lys	Asn	Val 1275	Leu	Asn	3956
atc c Ile L	Leu														4001

ı, °

							aag Lys 1300								4046		
							tgc Cys 1315								4091		
							cag Gln 1330								4136		
							ctg Leu 1345								4181		
							tgg Trp 1360						cac His	_	4226		
							cag Gln 1375								4271		
							gtg Val 1390								4316		
							gtg Val 1405						acc Thr		4361	÷	
							atg Met 1420								4406		
							ttc Phe 1435							ctg Leu	4451		
							aag Lys 1450							acg Thr	4496		
						Pro	ccc Pro 1465								4541		
							gtg Val 1480								4586		
							atg Met 1495						atc Ile	~ ~	4631		
ttc	acg	tcc	atg	ttc	tcc	atg	gag	tgt	gtg	ctc ·	aag	gtc	atc	gcc	4676		•

Phe	Thr	Ser 1505	Met	Phe	Ser	Met	Glu 1510	Cys	Val	Leu	Lys	Val 1515	Ile	Ala			
							aga Arg 1525								4721		
	_			_		-	att Ile 1540		_			_	aca Thr		4766		
							ctc Leu 1555								4811		
							ctc Leu 1570								4856		
						Val	cag Gln 1585								4901	-	
							ctg Leu 1600								4946		
							att Ile 1615								4991		
							cgg Arg 1630								5036		
							999 Gly 1645								5081		
_		_		-	_		gcc Ala 1660	_	_			_		-	5126		
							gcc Ala 1675								5171		
							atg Met 1690								5216		
							ctt Leu 1705								5261		
				_	_	_	ttc Phe			_		-			5306		

	1715					1720					1725				
gac ccg Asp Pro														5351	
atg ctg Met Leu														5396	
cct gct Pro Ala														5441	
atc tcc Ile Ser														5486	
gcc ctc Ala Leu														5531	
acg aag Thr Lys	cag Gln 1805	cat His	cag Gln	tgt Cys	gac Asp	gca Ala 1810	gag Glu	ctg Leu	agg Arg	aag Lys	gag Glu 1815	atc Ile	tcc Ser	5576	
tcc gtg Ser Val														5621	
cca ccc Pro Pro														5666	£-
gct ctg Ala Leu	-										acc Thr 1860		aga Arg	5711	
gac cag Asp Gln														5756	
gtg tcc Val Ser														5801	
ccg gct Pro Ala														5846	
tcg gcc Ser Ala														5891	
ggc atc Gly Ile														5936	

· •

_	_		_				_		_		cac His 1950			5981		-
	_						Pro	_		_	gtc Val 1965	_	_	6026		
											caa Gln 1980			6071		
_	 _	_		_	_	_		_		_	ctg Leu 1995		•	6116		
											cgc Arg 2010			6161		
	-	_		_					_		ctg Leu 2025	_		6206		
											ccc Pro 2040			6251	•	
	_	_		_	_		_		_	_	aag Lys 2055		•	6296		
				_		_	_		_		gcg Ala 2070			6341		
											gat Asp 2085			6386		
_	 _				_	agg Arg 2095	_			-	cgg Arg 2100		cag Gln	6431		
	 	_									cag Gln 2115	_		6476		
		Asp					Arg				cag Gln 2130			6521		
											999 Gly 2145	_		6566		

		ccc Pro 2150	Pro				agc Ser 2155							ccc Pro	6611
		tcg Ser 2165					agc Ser 2170					ggc Gly 2175		cgg Arg	6656
	_			_		_	ctg Leu 2185			_		agc Ser 2190		acc Thr	6701
	_	_	_				cct Pro 2200	_			_		_	_	6746
	_			_			ccc Pro 2215				_	_		ctt Leu	6791
_	_	cac His 2225		_	_		cag Gln 2230	_	_			_	_	_	6836
		ccc Pro 2240					ggc Gly 2245							cag Gln	6881
_	_	_	_		_		gcc Ala 2260	_		_			_	acg Thr	6926
		ttc Phe 2270				-	gcc Ala 2275			_		cgc Arg 2280		tcc Ser	6971
		tcc Ser 2285					ctg Leu 2290		_	_		cac His 2295		ctc Leu	7016
		gtg Val 2300					cac His 2305							tcg Ser	7061
							cac His 2320	_						caa Gln	7106
_		tgg Trp 2330	_	tag											7121

<210> 44 <211> 2331 <212> PRT <213> Bos taurus

- Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly 1 5 10 15
- Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Val Ala Gly Gly Pro 20 25 30
- Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser 35 40 45
- Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val 50 60
- Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu 65 70 75 80
- Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro 85 90 95
- Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu 100 105 110
- Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu 115 120 125
- Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu 130 135 140
- Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Leu His Lys Gly Ser 145 150 155 160
- Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr 165 170 175
- Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg 180 \$180\$
- Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser 195 200 205
- Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu 210 215 220

Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn 245 250 Ser Thr Asp Ala Asp Pro Val Gly Asp Phe Pro Cys Gly Arg Glu Ala 265 Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Ala Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile 295 Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu 310 315 Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe 330 Leu Pro Leu Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu 340 345 Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn 355 360 365 Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu 370 375 Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu 385 Ala Glu Glu Asp Lys Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu

Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu

Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe

440

Gln Ile Gly Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile

230

405

420

435

Ala	Arg 450	Ala	Ser	Leu	Lys	Ser 455	Gly	Lys	Thr	Glu	Ser 460	Ser	Ser	Tyr	Phe
Arg 465	Arg	Lys	Glu	Lys	Met 470	Phe	Arg	Phe	Phe	Ile 475	Arg	Arg	Leu	Val	Lys 480
Ala	Gln	Ser	Phe	Tyr 485	Trp	Val	Val	Leu	Cys 490	Val	Val	Ala	Leu	Asn 495	Thr

Leu Cys Val Ala Met Val His Tyr Gln Gln Pro Gln Arg Leu Thr Thr 500 505 510

Ala Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu 515 520 525

Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser 530 535 540

Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Ile Phe Glu 545 550 555 560

Val Val Trp Ala Ala Ile Lys Pro Gly Thr Ser Phe Gly Ile Ser Val 565 570 575

Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp 580 595

Ser Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser 595 600 605

Ile Ile Ser Leu Leu Phe Leu Phe Leu Phe Ile Val Val Phe Ala 610 615 620

Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu 625 630 635 640

Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val 645 650 655

Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly 660 665 570

- Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Val Tyr 675 680 685
- Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe 690 695 700
- Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys
  705 710 715 720
- Asp Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln
  725 730 735
- Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile 740 745 750
- Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp 755 760 765
- Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys 770 780
- Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala 785 790 795 800
- Thr Ser Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro 805 810 815
- Leu Val Val Glu Pro Gly Arg Asp Gly Ala Arg Gly Pro Ala Gly Gly 820 825 830
- Lys Ala Arg Pro Glu Gly Glu Glu Glu Glu Gly Ala Asp Pro Pro 835 840 845
- Arg Arg His His Arg His Arg Asp Arg Asp Arg Asp Lys Ala Ala 850 860
- Pro Ala Gly Glu Pro Asp Arg Ala Asp Ala Pro Lys Ala Glu Gly Gly 865 870 875 880
- Glu Pro Gly Ala Arg Glu Glu Arg Ala Arg Pro Arg Arg Ser Arg Ser 885 890 895
- Lys Glu Ala Ala Gly Pro Arg Glu Ala Arg Ser Glu Arg Gly Arg Gly

900 905 910

Leu Gly Pro Asp Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu 915 920 925

- Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Ala 930 935 940
- Pro Glu Pro Ser Arg Glu Gly Ala Pro Gly Ser Lys Gly Glu Arg Arg 945 950 955 960
- Ala Arg His Arg Gly Gly Ser Arg Ala Gly Pro Arg Glu Ala Glu Ser 965 970 975
- Gly Glu Glu Pro Ser Arg Arg His Arg Ala Arg His Lys Ala Pro Pro 980 985 990
- Ala His Glu Glu Ala Glu Lys Glu Ala Glu Ala Glu Asp Arg Asp Lys 995 1000 1005
- Glu Pro Arg Asn His Gln Pro Arg Glu Ser His Cys Asp Leu Glu 1010 1015 1020
- Ala Ser Val Val Ile Gly Val Gly Pro Val His Ala Leu Pro Ser 1025 1030 1035
- Thr Cys Leu Glu Lys Val Glu Glu Gln Pro Glu Asp Ala Asp Asn 1040 1045 1050
- Gln Arg Asn Val Thr Arg Met Gly Ser Gln Pro Ser Asp Leu Ser 1055 1060 1065
- Thr Ala Val Arg Ile Pro Val Thr Leu Thr Gly Pro Pro Gly Glu 1070 1075 1080
- Thr Thr Val Val Pro Ser Gly Asn Val Asp Leu Glu Ser Gln Ala 1085 1090 1095
- Glu Gly Lys Lys Glu Val Glu Ala Asp Asp Val Met Arg Ser Gly 1100 1105 1110
- Pro Arg Pro Ile Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro 1115 1120 1125

- Thr Asn Leu Leu Arg Arg Phe Cys His Tyr Ile Val Thr Met Gln 1130 1135 1140
- Tyr Phe Glu Met Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile 1145 1150 1155
- Ala Leu Ala Ala Glu Asp Pro Val Gln Thr Asp Ser Pro Arg Asn 1160 1165 1170
- Asn Val Leu Lys Tyr Met Asp Tyr Ile Phe Thr Gly Val Phe Thr 1175 1180 1185
- Phe Glu Met Val Ile Lys Met Ile Asp Leu Gly Leu Leu His 1190 1195 1200
- Pro Gly Ala Tyr Phe Arg Asp Leu Trp Asn Ile Leu Asp Phe Ile 1205 1210 1215
- Val Val Ser Gly Ala Leu Val Ala Phe Ala Phe Ser Gly Ser Lys 1220 1225 1230
- Gly Lys Asp Ile Ser Thr Ile Lys Ser Leu Arg Val Leu Arg Val 1235 1240 1245
- Leu Arg Pro Leu Lys Thr Ile Lys Arg Leu Pro Lys Leu Lys Ala 1250 1255 1260
- Val Phe Asp Cys Val Val Asn Ser Leu Lys Asn Val Leu Asn Ile 1265 1270 1275
- Leu Ile Val Tyr Met Leu Phe Met Phe Ile Phe Ala Val Ile Ala 1280 1285 1290
- Val Gln Leu Phe Lys Gly Lys Phe Phe Tyr Cys Thr Asp Glu Ser 1295 1300 1305
- Lys Glu Leu Glu Arg Asp Cys Arg Gly Gln Tyr Leu Asp Tyr Glu 1310 1315 1320
- Lys Glu Glu Val Glu Ala Gln Pro Arg Gln Trp Lys Lys Tyr Asp 1325 1330 1335

- Phe His Tyr Asp Asn Val Leu Trp Ala Leu Leu Thr Leu Phe Thr 1340 1345 1350
- Val Ser Thr Gly Glu Gly Trp Pro Met Val Leu Lys His Ser Val 1355 1360 1365
- Asp Ala Thr Tyr Glu Glu Gln Gly Pro Ser Pro Gly Phe Arg Met 1370 1380
- Glu Leu Ser Ile Phe Tyr Val Val Tyr Phe Val Val Phe Pro Phe 1385 1390 1395
- Phe Phe Val Asn Ile Phe Val Ala Leu Ile Ile Thr Phe Gln 1400 1405 1410
- Glu Gln. Gly Asp Lys Val Met Ser Glu Cys Ser Leu Glu Lys Asn 1415 1420 1425
- Glu Arg Ala Cys Ile Asp Phe Ala Ile Ser Ala Lys Pro Leu Thr 1430 1440
- Arg Tyr Met Pro Gln Asn Lys Gln Ser Phe Gln Tyr Lys Thr Trp 1445 1450 1455
- Thr Phe Val Val Ser Pro Pro Phe Glu Tyr Phe Ile Met Ala Met 1460 1465 1470
- Ile Ala Leu Asn Thr Val Val Leu Met Met Lys Phe Tyr Asp Ala 1475 1480 1485
- Pro Tyr Glu Tyr Glu Leu Met Leu Lys Cys Leu Asn Ile Val Phe 1490 1495 1500
- Thr Ser Met Phe Ser Met Glu Cys Val Leu Lys Val Ile Ala Phe 1505 1510 151.5
- Gly Val Leu Asn Tyr Phe Arg Asp Ala Trp Asn Val Phe Asp Phe 1520 1530
- Val Thr Val Leu Gly Ser Ile Thr Asp Ile Leu Val Thr Glu Ile 1535 1540 1545

- Ala Asn Asn Phe Ile Asn Leu Ser Phe Leu Arg Leu Phe Arg Ala 1550 1560
- Ala Arg Leu Ile Lys Leu Leu Arg Gln Gly Tyr Thr Ile Arg Ile 1565 1570 1575
- Leu Leu Trp Thr Phe Val Gln Ser Phe Lys Ala Leu Pro Tyr Val 1580 1585 1590
- Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile Gly 1595 1600
- Met Gln Val Phe Gly Asn Ile Ala Leu Asp Asp Asp Thr Ser Ile 1610 1615 1620
- Asn Arg His Asn Asn Phe Arg Thr Phe Leu Gln Ala Leu Met Leu 1625 1630 1635
- Leu Phe Arg Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu 1640 1650
- Ser Cys Leu Ser Ser Arg Ala Cys Asp Glu Leu Ala Asn Ala Thr 1655 · 1660 1665
- Glu Cys Gly Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile 1670 1680
- Phe Leu Cys Ser Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile 1685 1690 1695
- Met Asp Asn Phe Glu Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly 1700 1705 1710
- Pro His His Leu Asp Glu Phe Ile Arg Val Trp Ala Glu Tyr Asp 1715 1720 1725
- Pro Ala Ala Cys Gly Arg Ile Ser Tyr Asn Asp Met Phe Glu Met 1730 1740
- Leu Lys His Met Ser Pro Pro Leu Gly Leu Gly Lys Lys Cys Pro 1745 1750 1755
- Ala Arg Val Ala Tyr Lys Arg Leu Val Arg Met Asn Met Pro Ile

1760	1765	1770

Ser	Asn 1775	Asp	Asp	Met	Thr	Val 1780	His	Phe	Thr	Ser	Thr 1785	Leu	Met	Ala
Leu	Ile 1790	Arg	Thr	Ala	Leu	Glu 1795		Lys	Leu	Ala	Pro 1800	Ala	Gly	Thr
Lys	Gln 1805	His	Gln	Cys	Asp	Ala 1810	Glu	Leu	Arg	Lys	Glu 1815	Ile	Ser	Ser
Val	Trp 1820	Ala	Asn	Leu	Pro	Gln 1825	_	Thr	Leu	Asp	Leu 1830	Leu	Val	Pro
Pro	His 1835	Lys	Pro	Asp	Glu	Met 1840	Thr	Val	Gly	Lys	Val 1845	Tyr	Ala	Ala
Leu	Met 1850	Ile	Phe	Asp	Phe	Tyr 1855	Lys	Gln	Asn	Lys	Thr 1860	Thr	Arg	Asp
Gln	Ile 1865	His	Gln	Ala	Pro	Gly 1870	Gly	Leu	Ala	Gln	Met 1875	Gly	Pro	Val
Ser	Leu 1880	Phe	His	Pro	Leu	Lys 1885	Ala	Thr	Leu	Glu	Gln 1890	Thr	Gln	Pro
Ala	Val 1895	Leu	Arg	Gly	Ala	Arg 1900	Val	Phe	Leu	Arg	Gln 1905	Lys	Ser	Ser
Ala	Ser 1910	Leu	Ser	Asn	Gly	Gly 1915	Ala	Val	Gln	Thr	Gln 1920	Glu	Gly	Gly
Ile	Lys 1925	Glu	Ser	Val	Ser	Trp 1930	Gly	Thr	Gln	Arg	Thr 1935	Gln	Glu	Val
Pro	Cys 1940	Glu	Val	Arg	Thr	Pro 1945	Leu	Glu	Arg	Gly	His 1950	Ser	Thr	Glu
Ile	Pro 1955	Val	Gln	Thr	Gly	Lys 1960	Pro	Ala	Val	Asp	Val 1965	Gln	Met	Gln
Ser	Met 1970	Val	Leu	Arg	Gly	Pro 1975	Asp	Gly	Glu	Pro	Gln 1980	Pro	Gly	Leu

- Glu Ser Gln Gly Arg Ala Ala Ser Met Pro Arg Leu Ala Ala Glu 1985 1990 1995
- Thr Gln Pro Pro Pro Asp Ala Ser Pro Met Lys Arg Ser Ile Ser 2000 2005 2010
- Thr Leu Ala Pro Gln Arg Pro His Val Ala His Leu Cys Thr Ala 2015 2020 2025
- Ala Leu Asp Arg Ala Pro Ala Ser Gln Ala Ala Pro Pro His His 2030 2035 2040
- His Arg Cys His Arg Arg Arg Asp Arg Lys Gln Lys Ser Leu Glu 2045 2050 2055
- Lys Gly Pro Gly Leu Ser Ala Asp Thr Asp Gly Ala Pro Tyr Ser 2060 . 2065 2070
- Thr Ala Gly Pro Gly Pro Pro Pro Pro Gly Asp Gly Ala Ala 2075 2080 2085
- Gly Cys Arg Arg Glu Arg Arg Gln Glu Arg Gly Arg Ser Gln Glu 2090 2095 2100
- Arg Arg Gln Pro Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr 2105 2110 2115

. 3

- Ser Cys Asp Arg Phe Gly Gly Arg Glu Pro Pro Gln Pro Lys Pro 2120 2125 2130
- Ser Leu Ser Ser His Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro 2135 2140 . 2145
- Gly Pro Pro Arg Gln Gly Ser Gly Ser Val Asn Gly Ser Pro Leu 2150 2155 2160
- Leu Ser Thr Ser Gly Ala Ser Thr Pro Gly Arg Gly Gly Arg Arg 2165 2170 2175
- Gln Leu Pro Gln Thr Pro Leu Thr Pro Arg Pro Ser Ile Thr Tyr 2180 2185 2190

Lys Thr Ala Asn Ser Ser Pro Val His Phe Ala Gly Ala Gln Thr 2195 2200 2205

Ser Leu Pro Ala Phe Ser Pro Gly Arg Leu Ser Arg Gly Leu Ser 2210 2215 2220

Glu His Asn Ala Leu Leu Gln Arg Asp Pro Leu Ser Gln Pro Leu 2225 2230 2235

Ala Pro Ser Ser Arg Ile Gly Ser Asp Pro Tyr Leu Gly Gln Arg 2240 2245 2250

Leu Asp Ser Glu Ala Ala Ala Arg Thr Gln Leu Glu Asp Thr Leu 2255 2260 2265

Thr Phe Glu Glu Ala Val Ala Thr Asn Ser Gly Arg Ser Ser Arg 2270 2275 2280

Thr Ser Tyr Val Ser Ser Leu Thr Ser Gln Ser His Pro Leu Arg 2285 2290 2295

Arg Val Pro Asn Gly Tyr His Cys Thr Leu Gly Leu Ser Ser Gly 2300 2310

Gly Gly Arg Gly Arg His Ser Tyr His His Pro Asp Gln Asp 2315 2320 2325

His Trp Cys 2330

<210> 45

<211> 33

<212> PRT

<213> Homo sapiens

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Amino acid sequence encoded by exon 37a; amino acid at position 1 is partly encoded by exon 36, partly by exon 37a. Identical amino acid sequence is found in Homo sapiens, Mus musculus and Rattus norvegicus

<400> 45

Cys Cys Arg Ile His Tyr Lys Asp Met Tyr Ser Leu Leu Arg Cys Ile 1 5 10 15

Ala Pro Pro Val Gly Leu Gly Lys Asn Cys Pro Arg Arg Leu Ala Tyr 20 25 30

Lys

<210> 46

<211> 32

<212> PRT

<213> Rattus norvegicus

<400> 46

Cys Cys Arg Ile His Tyr Lys Asp Met Tyr Ser Leu Leu Arg Cys Ile 1 5 10 15

Ala Pro Pro Val Gly Leu Gly Lys Asn Cys Pro Arg Arg Leu Ala Tyr
20 25 30

<210> 47

<211> 32

<212> PRT

<213> Rattus norvegicus

<400> 47

Cys Gly Arg Ile Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His Met 1 5 10 15

Ser Pro Pro Leu Gly Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr 20 25 30